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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                      Score
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length: 12
                                                                                                                                                                                                                                                                                                                     Query
Match
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112.614 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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59
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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30.5
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A58718
$78420
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B46453
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828055
8137268
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857873
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SPPGNK
C61033
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A55837
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S35490
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NADH2 dehydronenn
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alpha-myosin heavy
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neuromedin K - pig
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tetracenomycin A2
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type II site-speci
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pilE protein - Esc
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                                                                                                       glycoprotein H-a
                                                                                                                                                                    polyphosphate-gluc
chlorophyll a/b-bi
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                                                                                              -aminoimidazole r
proline-ric
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$21205

Ig heavy chain V region - human

C;Species: Homo sapiens (man)

C;Species: Homo sapiens (man)

C;Species: 2-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 16-Aug-1996

C;Accession: $21205

R;Makiya, R.; Stigbrand, T.

Bur. J. Biochem. 205, 341-345, 1992

Bur. J. Biochem. 205, 341-345, 1992

A;Title: Placental alkaline phosphatase has a binding site for the human immunoglobulin-
A;Reference number: $21205; MUID:92209522; PMID:1555592

A;Accession: $21205
                                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-12 «MAK»
C;Keywords: heterotetramer; immunoglobulin
  RESULT
S78420
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                                            SEIQ 5
                                                                 SEIQ 6
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ALIGNMENTS

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C;Accession: A58718
R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, I.F. Appl. Environ. Microbiol. 58, 1417-1422, 1992
A;Title: Purification and characterization of a new bacteriocin isolated from a Carnoba A;Reference number: A58718; MUID:92321768; PMID:1622206
A;Accession: A58718
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 <STO-
                                                                                                                                                                                                                                                                                                                                                                   carnocin UI49 - Carnobacterium sp. (fragment)
C;Species: Carnobacterium sp.
C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
                                                                                                        A;Residues: 1-7 <STO>
C;Keywords: antibiotic; lanthionine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 35.6%; Score 21; DB 2; Le Similarity 44.4%; Pred. No. 6.5e+02; 4; Conservative 3; Mismatches 2;
30.5%; Score 18; DB 2; Li ilarity 100.0%; Pred. No. 2.8e+05; Conservative 0; Mismatches 0;
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sperm-activating peptide (Tyr-2, Asn-3, Gly-5, Ile-9, Asp-10 SAP-I) - slate-pencil urchi (JSpecies: Heterocentrotus mamillatus (JDate: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000 C;Accession: A60589
C;Accession: A60589
C;Accession: A60589
C;Accession: Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, M.; Yamaguc Comp. Blochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and its related peotus mudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Accession: A60589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Accession: S78420
R;Goldschmidt-Reisin, S.; Graack, H.R.
submitted to the Protein Sequence Database, February 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ribosomal protein RL41, mitochondrial [validated] - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb_1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                   C;Date: 18-Jun-1993 #sequence_revision 08-Nov-1996 #text_change 15-Aug-1997 C;Accession: B46453  
R;Takahashi, K.; Kishimoto, S.; Ohori, K.; Yoshizawa, H.; Machida, A.; Ohnuma, H. J. Immunol. 147, 3156-3160, 1991  
A;Title: Molecular heterogeneity of e antigen polypeptides in sera from carriers A;Reference number: A46453; MUID:92013147; PMID:1717588
                                                                                                                                                                                                                                                                                                                                                          e antigen p20e precursor - hepatitis B virus N;Alternate names: HBe antigen precursor N;Contains: e antigen C;Species: hepatitis B virus, HBV A;Variety: subtype adr C;Date: 1B = Jun-1993 #sequence_revision 08-Nov C;Accession: B46453
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Note: the protein is designated a C;Keywords: mitochondrion; protein
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A; Accession: S78420
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Best Local S
Matches 3
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Matches 3
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Experimental source: subtype adr

Y-1--- convence extracted from NCBI backbone (NCBIP:60243)
                                                                                                                                                                                          Molecule type: protein
Residues: 1-10 <TAK>
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Pred. No. 2e+03;
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Pred. No. 2.8e+05;
                                                                  Score 18; DB 2;
Pred. No. 2e+03;
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C;Accession: I46868
R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984
A;Title: (haracterization of genomic clones specifying A;Reference number: I46868; MUID:84221901; PMID:6328491
A;Accession: I46868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: The psbL gene from A;Reference number: S28055; A;Accession: S28055
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R;Kuntz, M.; Camara, B.; Weil, J.H.; Schantz,
Plant Mol. Biol. 20, 1185-1188, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytochrome b559 component psbr - pepper chloroplast (fragment)
C;Species: chloroplast Capsicum annuum (pepper)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_chang
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S28055
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A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-10 <RUF
                                                                                            R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R
J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences
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C;Keywords: chloroplast; ]
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A;Residues: 1-10 <KUN>
A;Cross-references: EMBL:X65570;
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C;Sate: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change
                                                      A;Reference number: A38740; MUID:91177923; PMID:1706720
                                                                                                                                                     Ig heavy chain C region (129) - mouse (C;Species: Mus musculus (house mouse) C;Date: 19-Mar-1997 #sequence_revision C;Accession: A37268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Genome: chloroplast
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                                                                                                                                                                                                                  mouse (fragment)
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Pred. No. 2.8e+05;
1; Mismatches 1
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Pred. No. 3.1e+03
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                                                                                                     antibody properties
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A; Molecule type: protein A; Residues: 1-9 <SER>
C; Superfamily: co---
C; Keywor--
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N;Alternate names: beta-COP protein
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 19-Mar-1997 #sequence_revision 24-Mar-1999 #text_chan
C;Accession: S13636
R;Serafini, T.; Stenbeck, G.; Brecht, A.; Lottspeich, F.; Orc
Nature 349, 215-220, 1991
A;Title: A coat subunit of Golgi-derived non-clathrin-coated
A;Reference number: S13636; MUID:91101693; PMID:1898984
A;Accession: S13636
                                   pilE protein - Escherichia coli (fragment)
C;Species: Escherichia coli
C;Species: Bscherichia coli
C;Date: 28-May-193 #sequence_revision 28-May-1993 #text_change 11-Jun-1993
C;Accession: $25266
C;Accession: $25266
Mol. Microbiol. 6, 1887-1894, 1992
Mol. Microbiol. 6, 1887-1894, 1992
A;Fitle: Pulo, a component of the pullulanase secretion pathway of Klebsiel
A;Reference number: $25266; MUID:92374839; PMID:1354833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            placental lactogen-I precursor - mouse (fragment)
C;Species: Mus sp. (mouse)
C;Species: Nus sp. (mouse)
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996
C;Accession: A47364
R;Shida, M.M.; Ng, Y.K.; Soares, M.J.; Linzer, D.I.
Mol. Endocrinol. 7, 181-188, 1993
A;Title: Trophoblast-specific transcription from the mouse placental lactoge A;Accession: A47364
A;Accession: A47364
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A47364
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A;Accession: S25266
A;Molecule type: prot
A;Residues: 1-7 <DUP>
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Best Local S
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Best Local
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57.1%;
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Pred. No. 2.8e
3; Mismatches
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Pred. No. 3.1e+03;
1; Mismatches 4;
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2.8e+05;
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PW0002
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                                                 N-methylpurine DNA glycosylase - mouse (fragment) (;Species: Mus musculits (house mouse) (;Date: 19-Jan-196 #sequence_revision 19-Jan-199 C;Accession: A56029
R;Roy, R.; Brooks, C.; Mitra, S.
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 R;Roy, R.; Brooks, C.; Mitra, S.
Biochemistry 33, 15131-15140, 194
A;Title: Purification and biochemical characterization of recombinant N-methylpurine-DNA
A;Reference number: A56029; MUID:95092772; PMID:7999773
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3; Conserv
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polyphosphate-glucose phosphotransferase (EC 2.7.1.63) - Propionibacterium freudenreichi C,Species: Propionibacterium freudenreichi subsp. shermanii C,Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 03-Jun-2002 C;Accession: S29735 #sequence_revision 19-Mar-1997 #text_change 03-Jun-2002 R;Phillips, N.F.B.; Horn, P.J.; Wood, H.G. Arch. Biochem. Biophys. 300, 309-319, 1993 A;Title: The polyphosphate- and ATP-dependent glucokinase from Propionibacterium sherman A;Reference number: S29735; MUID:93143332; PMID:8380966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 eHI>
C;Keywords: phosphotransferase
                                                                                                                                                                                      R:LaRoche, J.; Bennett, J.; Falkowski, P.G.
Gene 95, 165-171, 1990
A;Title: Characterization of a cDNA encoding for the 28.5-kDa
A;Reference number: JW0040; MUID:91065528; PMID:2249775
                                                                                                                                                                                                                                                                                             chlorophyll a/b-binding protein 31% - green alga (D N;Alternate names: photosystem II light-harvesting C;Species: Dunalisella tertiolecta C;Date: 24-Nov-1999 #sequence_revision 24-Nov-1999
                                                                                    C; Superfamily: chlorophyll a/b-binding protein C; Keywords: chloroplast; grana; light-harvesting
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                      Score 15;
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Pred. No. 2.8e+05;
1; Mismatches 0;
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Pred. No. 2.8e+0
0; Mismatches
        Mismatches
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..8e+05;
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ng chlorophyll 31K protein
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19-Jan-1996

#text_change 19-Jan-1996

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hypothetical protein 2 LRH-1 5'-region - mouse
C:Species: Mus musculus (house mouse)
C:Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
C:Accession: S27873
R;Tugwood, J.D.; Issemann, I.; Green, S.
submitted to the EMBL Data Library, February 1992
A:Description: LRH-1: A nuclear hormone receptor active in the absence of exogenous ligal A;Reference number: S27873
A;Reference number: S27873
A;Accession: S27873
A;Accession: S27873
A;Residues: 1-10 cTUG>
A;Cross-references: EMBL:M81385; NID:g198872; PIDN:AAA39446.1; PID:g198874
C;Superfamily: unassigned leader peptides
Search completed: May 18, 2004, 10:05:40 Job time: 11.25 secs
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-9 <ROY>
                                                                                                                                                                                                    Query Match 25.4%; Score 15; DB 2; Length 10; Best Local Similarity 50.0%; Pred. No. 7.6e+03; Matches 3; Conservative 2; Mismatches 1; Indels
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                                                                                               5 NLSHIQ 10
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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ALLI CRAMA
ASHI BACSE
CD14 LITXA
CC14 LITXA
CC14 LITXA
CC14 LEUMA
ACSI BACSU
PRIA PERRM
CS15 BACSU
Q20A COMTE
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PRIA PERRM
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RESULT 2

PSBF CAPAN

ID PSBF CAPAN

AC Q03367;
DT 01-JUN-1994 (Rel. 29, Created)
DT 01-JUN-1994 (Rel. 29, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Cytochrome b559 beta subunit (PSII reaction cent. Fragment).

GN PSBF.

OS Capsicum annuum (Bell pepper).

CC Chloroplast.

OC Eukaryota; Viridiplantae; Streptophyta; Embryoph;
OC Eukaryota; Viridiplantae; Streptophyta; Embryoph;
OC Spermatophyta; Magnoliophyta; eudicotyledons; co:
OC 1amiids; Solanales; Solanaceae; Capsicum.

CX NCBI_TaxID=4072;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=cv. Lammyo; TISSUE=Fruit, and Leaf;
RX MEDLINE=93099270; PubMed=1463853;
RA Kuntz M., Camara B., Weil J.-H., Schantz R.;
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ID LANC_C
AC P36960
DT 01-UN
DT 16-OCT
DE LantLb
OS Carnob
OC Bacter
OC Carnob
OC MCBI_T
RN [1]
RN [1]
RN SLOLIN
RY MEDLLIN
RX MEDLL
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; aster lamiids; Solanales; Solanaceae; Capsicum.
NCBI_TaxID=4072;
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P36960;
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NCBI_TaxID=35782;
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ALIGNMENTS	PIBB_PAPHA RE42_LITRU GLEM_HUWAN GON1_CLUPA GON2_CHEPR HTF1_ROWMI HTF_HELZE SYK_CAMUP TKNB_RANRI CXL1_COMMR HS70_PINPS HEP1_BACSE
	P19343 papio hamad P82075 litoria rub P82075 litoria rub P02728 homo sapien P81749 clupea pall P816749 chelyosoma p18110 romalea mic P16353 heliothis z Q46464 campylobact P29135 rana ridibu P58807 conus marmo P58807 opius pinas P83054 bacteroides

a Carnobacterium sp."; Appl. Environ. Microbiol. 58:1417-1422(1992). -i- FUNCTION: Lanthionine-containing peptide Active on Gram-positive bacteria. Antibiotic; Bacteriocin; Lantibiotic. Carnobacterium sp. (strain UI49). Bacteria; Firmicutes; Lactobacillales; Carnobacteriaceae; 01-JUN-1994 (Rel. 29, Created) 01-JUN-1994 (Rel. 29, Last sequence up 16-CCT-2001 (Rel. 40, Last annotation Lantibiotic carnocin UI49 (Fragment). Carnobacterium sp. (strain UI49). MEDINE=92321768; PubMed=1622206; Stoffels G., Nissen-Meyer J., Gud "Purification and characterization of a new bacteriocin isolated from 7 AA; 786 MW; 30.5%; pre 100.0%; pre 741776D05B05B810 CRC64; Score 18; DB 1; Pred. No. 1.4e+05; Gudmundsdottir A., PRT; on update) Ā antibiotic (lantibiotic) Length 7; Sletten Indels ζ. 0, Holo Gaps 0

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P82157;
30-MAY-2000 (Rel. :
30-MAY-2000 (Rel. :
30-MAY-2000 (Rel. :
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                                                                                                                                                                                                                                                                                                                                   Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstan Davey M., East P.D., Thorpe A.;
"Lepidopreran peptides of the allatostatin superfamily.";
Peptides 18:1301-1309(1997).
-!- SIMILARITY: Belongs to the allatostatin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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Cydia pomonella (Codling moth)
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Chloroplast; Photosystem II; Heme; Electron transport; Transmembrane.
NON TER
1 1
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BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "The psb1 gene from bell pepper (Capsicum annuum): plastid RNA editing also occurs in non-photosynthetic chromoplasts."; Plant Mol. Biol. 20:1185-1188 [1992].
-!- FUNCTION: This b-type cytochrome is tightly associated with the reaction center of photosystem II and possibly is part of the water-oxidation complex.
-!- SUBUNIT: Heterodimer of an alpha subunit and a beta subunit.
-!- SIMILARITY: Belongs to the psb5 / psbF family.
                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                            Neuropeptide; Amidation.
MOD_RES 8 8
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MEDLINE=98054539; PubMed=9392829;
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InterPro; IPR006216; Cyt_b559.
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37.5%;
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0B2879C45B573767 CRC64;
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LUMENAL (POTENTIAL).
817D0F59D6D69DC5 CRC64;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scott A.G., Winstanley D.
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P35946;
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Submitted (JUL-1997)
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                              Transferase; RNA-directed RNA
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                                                                                 EMBL; D10371; BAA01208.1
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RESULT

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1105 MW;

9C2B7FD452D5A2D5 CRC64

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psiolo;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 42, Last annotation update)
10-OCT-2003 (Rel. 42, Last annotation update)
Allergen Fus s 13596* (Fragment).
Fusarium solani (subsp. pisi) (Nectria haematocc Eukaryota; Fungi; Ascomycota, Pezizomycotina; Sc Hypocreomycetidae; Hypocreales; Nectriaceae; Nec NCBI_TaxID=70791;
                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=Ulster/88;

MEDLINE=92268977; PubMed=1588321;

Curran M.D., O'Loan D., Kennedy S., Rima B.K.;

Curran M.D., O'Loan D., Kennedy S., Rima B.K.;

"Molecular characterization of phocine distemper virus: gene order and sequence of the gene encoding the attachment (H) protein.";

and sequence of the gene encoding the active polymerase. It may J. Gen. Virol. 73:1189-1194(1992).

-!- FUNCTION: Probable component of the active polymerase. It may function in mRNA synthesis, capping, methylation and poly(A) synthesis of newly synthesized viral mRNAs, RNA editing of the P gene transcript, and protein kinase activity.

-!- CATALYTIC ACTIVITY: N nucleoside triphosphate = N diphosphate +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
RNA polymerase beta subunit (EC 2.7.7.48) (Last protein) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phocine distemper virus (PDV).
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.

    -!- SIMILARITY: Belongs to the paramyxoviruses L protein family.

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Causes an allergic reaction in human.
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polymerase
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RESULT 6
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1 PVKI PERAM STANDARD; PRT; 11 AA.

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InterPro, IPR000795, EF GTPbind.
PROSITE, PS00301; EFACTOR GTP; PARTIAL.
Blongarion factor; Protein biosynthesis,
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P81350;
              SEQUENCE, AND SYNTHESIS.
TISSUE-Abdominal perisympathetic organs;
MEDIINE-95212021; pubMed=7716075;
MEDIINE-95212021; pubMed=7716075;
Perdel R., Linde D., Rapus J., Vettermann S., Penzlin H.;
"Periviscerokinin (Pea-PVK): a novel myotropic neuropeptide
perisympathetic organs of the American cockroach.";
Peptides 16:61-66(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5. Electrophoresis 19:802-806(1998).

-i- FUNCTION: This protein promotes the GTP-dependent translocation protein chain from the A-site to the P-site of
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MEDLINE=98291870; PubMed=9629918;
Flengsrud R., Skjeldal L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JUL-1998 (Rel
15-JUL-1998 (Rel
28-FEB-2003 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Clostridium.
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Bacteria; Firmicutes; Cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Elongation factor
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. 36, Last
. 41, Last
c G (EF-G)
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25.0%;
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PEPTIDE;
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Pred. No. 3e+03;
2; Mismatches
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 HAS EXCITORY ACTIONS
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Matches 3
                01-JUN-1994
01-JUN-1994
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                                                                  PSEPU
TRPI PSEPU
P36414;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Cloning and characterization of genes for the PvuI restriction and modification system.",
Nucleic Acids Res. 20:5743-5747(1992).
-!- FUNCTION: RECOGNIZES THE DOUBLE-STRANDED SEQUENCE CGATCG AND CLEAVES AFTER T-4.
-!- CATALYRIC ACTUVITY: Endomucleolytic cleavage of DNA to give specific double-stranded fragments with terminal 5'-phosphates.
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P31031;
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PIR; S35490; S35490.
REBASE; 1541; PVuI.
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STRALN=ATCC 13315;
MEDLINE=93087186; PubMed=1454536;
Smith M.D., Longo M., Gerard G.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
10-CCT-2003 (Rel. 42, Last annotation update)
Type II restriction enzyme PvuI (EC 3.1.21.4)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; Proteobacteria;
Enterobacteriaceae; Proteo
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   HTH-type
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994 (Rel. 29, Created)
994 (Rel. 29, Last sequence update)
904 (Rel. 43, Last annotation update)
004 (Rel. 43, Last annotation update)
transcriptional regulator trpl (TrpBA operon transcriptional
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; Pred. No. 5.1e.
1; Mismatches
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Pred. No. 5
                                                                                                                                                                                                                                                                                                                   9F0CDE7955B72B1A CRC64;
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P82158;
30-MAY-2000
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STRAIN-PPG1 C1S;
MEDILINE=89315826; PubMed=2503057;
Eberly L., Crawford I.P.;
"DNA sequence of the tryptophan synthase
                                                       -!- SIMILARITY: Belongs to the allatostatin
Neuropeptide; Amidation.
MOD_RES 7 7 AMIDATION.
                                                                                   Duve H., Johnsen A.H., Maestro J.-L., Scott Davey M., East P.D., Thorpe A., "Lepidopteran peptides of the allatostatin Peptides 18:1301-1309(1997).

-!- SIMILARITY: Belongs to the allatostatir
                                                                                                                                      MEDLINE=98054539; PubMed=9392829;
Duve H., Johnsen A.H., Maestro J.-L.,
                                                                                                                                                                                                                    Cydia pomonella (Codling moth).
Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                                      CYDPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NO
NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000847; HTH_LYSR.
PROSITE; PS50931; HTH_LYSR; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biochimie 71:521-531(1989).
                                                                                                                                                                      TISSUE=Larva;
                                                                                                                                                                                                                                                                                   Cydiastatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; X13299; CAA31660.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pseudomonas
Bacteria; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tryptophan biosynthesis; Transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activator)
TRPI.
                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                           NCBI_TaxID=82600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION: ACTIVATES THE EXPRESSION OF THE TRPBA GENES ENCODING THE TWO TRYPTOPHAN SYNTHASE SUBUNITS. IN THE ABSENCE OF THE INDUCER (INDOLEGLYCEROL PHOSPHATE), TRPI BINDS UPSTREAM OF THE TRPAB OPERON, OVERLAPPING ITS OWN PROMOTER REGION.
SIMILARITY: Contains 1 HTH lysr-type DNA-binding domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                        LMHNL 11
                                                                                                                                                                                                                                                                                                                                                                                                                             MAHDL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Proteobacteria; Gammaproteobacteria; Pseudomonadales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 AA;
                                                                                                                                                                                                                                                                                               (Rel. 39, Created)
(Rel. 39, Last sequence up
(Rel. 39, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                         AA;
                                                                                                                                                                                                                                                                                                                                                        STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pseudomonas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         683 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23.7%;
 23.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        თ
                                         MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         77672AA1EDD6F000 CRC64;
                                         672879CABB569350
 Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. 1.4
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14;
                                                                                                                                                                                                                                                                                               on update)
                                                                                                                                         Scott
  DB 1;
1.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                       B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genes
                                                                                                             superfamily.";
                                                                                                                                         A.G., Winstanley D.,
                                                                                   family
             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 6;
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RESULT 11
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Best Local S
Matches 2
                     Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TALL1 CYDPO
P82152;
30-MAY-2000
30-MAY-2000
30-MAY-2000
                                                                                                    MOD RES
SEQUENCE
                                                                                                                                                Duve H., Johnsen A.H., Maestro J.-L., Scott Davey M., East P.D., Thorpe A.; "Lepidopteran peptides of the allatostatin septides 18:1301-1309(1997).

-! SIMILARITY: Belongs to the allatostatin Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P81821;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                Cydia pomonella (Codling moth)
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neuropeptide; Amidation; Multigene family.
MOD_RES 8 AMIDATION (POT)
SEQUENCE 8 AA; 919 MW; C82879D5AB569AB5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Isolation and identification of multiple neuropeptides of th allatostatin superfamily in the shore crab Carcinus maenas.", Eur. J. Biochem. 250:727-734 (1997).

FUNCTION: May act as a neurotransmitter or neuromodulator is remained by the allatostatin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Carcinus maenas (Common shore crab) (Green crab)
Eukaryota; Metazoa; Arthropoda; Crustacea; Malaco
Eumalacostraca; Eucarida; Decapoda; Pleocyemata;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                     Tortricoidea, Tor
NCBI_TaxID=82600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Cerebral ganglion, and Thoracic ganglion, MEDLINE-98121193; PubMed-9461295;
                                                                                                                                                                                                                                                                                                                                    SEQUENCE.
TISSUE=Larva,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cydiastatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thorpe A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Duve H.,
                                                                                                                                Neuropeptide;
                                                                                                                                                                                                                                                                                                           MEDLINE=98054539; PubMed=9392829;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=6759;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   മ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            œ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Rel. 39, Created)
(Rel. 39, Last seq
(Rel. 39, Last ann
                                                                                                    AA;
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                                                                                                       934 MW;
                        23.7%;
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Last annotation update)
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                                                                                                    C82879C45B51F775 CRC64;
Score 14; DB
Pred. No. 1.4e
1; Mismatches
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Pred. No. 1.4e+05;
2; Mismatches 1;
                                                                                                                          AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J.-L.,
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                                                                                                                                                                                                                                                                                Scott A.G., Winstanley
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                                                 DB 1;
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                                                                                                                                                                                                                               superfamily.";
                                                                                                                                                                                 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CRC64;
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                                                   Length
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Matches

Similarity 50.0 2; Conservative

.4e+05

Indels

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YNFG

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ALD CACALLA REPORT OF THE PROPERTY OF THE PROP
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P83281;
C P83281;
T 28-FEB-2003 (Rel. 41, Created)
T 28-FEB-2003 (Rel. 41, Last sequence update)
T 28-FEB-2003 (Rel. 41, Last sequence update)
E FMRPamide-like neuropeptide FLD8 (VSHNNFLRF-amide).
S Macrobrachium rosenbergii (Giant fresh water prawn).
C Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
C Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Caridea;
C Palaemonoidea; Palaemonidae; Macrobrachium.
N NCBI TaxID-79674;
Query Match 23.7%; Sometime Best Local Similarity 100.0%; 1 Matches 3; Conservative 0;
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Best Local (
                                                                                                               NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UF02_MOUSE STANDARD; PRT; 9 AA.

P38640;

01-OCT-1994 (Rel. 30, Created)

01-OCT-1994 (Rel. 30, Last sequence update)

15-MAR-2004 (Rel. 43, Last annotation update)

Unknown protein from 2D-page of fibroblasts (P32) (Fragment).

Mus musculus (Mouse).
                                                                                                                                            SEQUENCE.
TISSUB-Fibroblast;
MEDLINE-95009907; PubMed-7523108;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
MCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neuropeptide; Amidation.
MOD_RES 9 9
SEQUENCE 9 AA; 1133 M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Three more novel FMRFamide-like neuropeptide sequences from the eyestalk of the giant freshwater prawn Macrobrachium rosenbergii.", Peptides 22:191-197(2001), -i- MASS SPECTROMETRY: NMW-1133.8; METHOD-MALDI.
-i- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sithigorngul P., Saraithongkum W., Longyant S., Panchan N., Sithigorngul W., Petsom A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=21107394; PubMed=11179812;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE, AND MASS SPECTROMETRY.
TISSUE=Eyestalk;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            family.
GO:0007218; P:neuropeptide signaling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HN 10
                                                                                                               9 AA;
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                                                                                                                 1102 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1133 MW;
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                                                                                                                 7E73EAB6D05B1AAB CRC64;
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845A0729C44441F5
  Score 14; DB 1; Le
Pred. No. 1.4e+05;
0; Mismatches 0;
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Pred. No. 1.4e+
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Search completed: May 18, Job time: 7.75 secs

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Matches 2
                                                                                                                                                                                                                                                                                Liacostatin superfamily in the shore crab Carcinus maenas.",
Eur. J. Biochem. 250:727-734(197).

-!- FUNCTION: May act as a neurotransmitter or neuromodulator.

-!- SINLLARITY: Belongs to the allatostatin family.

Neuropeptide; Amidation; Multigene family.

MOD_RES

10
AMIDATION:
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Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu:
Eubrachyura; Portunoidea; Portunidae; Carcinus.
NCBI_TaxID=6759;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Cerebral ganglion, and Thoracic ganglion MEDLINE=98121193; PubMed=9461295;
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30-MAY-2000 (Rel
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                                                                                                                                                      Conservative
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. 39, Last sequence . 39, Last annotations.
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Pred. No. 7.2e+03;
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P82081 limnodynast
P82082 limnodynast
P82083 limnodynast
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O34919 locusta mig
O92766 cantine dist
O40659 oryza sativ
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O23876 dictyosteli
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O35891 homo sapien
O36012 linospadix
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ALIGNMENTS

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Matches 5; Conserv
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01-MAY-2000
01-MAY-2000
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Q9ES71;

Q1-MAY-2000 (TrEMBLrel. 13, Created)

Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)

Q1-MAY-2002 (TrEMBLrel. 21, Last annotation update)

Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)

Fibrinolytic metalloproteinase (Fragment).

Agkistrodon contortrix.

Eukaryota, Metazoa, Chordata, Craniata, Vertebrata; Euteleostomi;

Lepidosauria, Squamata, Scleroglossa, Serpentes, Colubroidea,

Wiperidae; Crotalinae; Agkistrodon.

NCBL_TaxID=8720;
                                                                                                                                                                                                                                                                                                                                                                                                           Guan A.L., Retzios A.D., Henderson G.N., Markland F.S.Jr.;
"Purification and characterization of a fibrinolytic enzyme from venom of the southern copperhead snake (Agkistrodon contortrix contortrix).";
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MEDLINE=91378546; PubMed=1898066;
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"ER 11 11
"NCE 11 AA; 1209 MW; 7CA02D1D41E87
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Pred. No. 1.4e+02;
1; Mismatches 1; Indels
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DYNASTIN 5.
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DYNASTIN 4.
Limnodynastes salmini (Salmin's-striped fr
Eukaryota; Metazoa; Chordata; Craniata; Ve
Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
NCBI TaxID=39404;
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-!- MASS
SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;

Limnodynastinae; Limnodynastes.
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TISSUE-TIBIAL GLAND;

Raftery M.J., Bradford A.M., Bowie J.H., Wallace J.C., Tyler M.J.;

"Peptides from Australian frogs. The structure of the dynastins from "Peptides from Australian frogs. The structure of the dynastins from the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii and the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., "Peptides from Australian frogs. The structure of the Limmodynastes salmini and Fletcherin from Limnodynaste Aust. J. Chem. 46:1235-1244(1993).
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Bukaryota; Metazoa; Chordata; Craniata; Va
Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
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Aust. J. Chem. 46:833-842(199)
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DYNASTIN 3.
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8 AA; 772 MW; 7B58772455A2C728 CRC64;
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SPECTROMETRY: MW=1236; METHOD=FAB.
12 AA; 1236 MW; 147AA70FDF472724
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(TrEMBLrel. 13, Last ann
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Pred. No. 1e+03;
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Pufonoidea; Myobatrachidae;
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Matches
SEQUENCE FROM N.A. MEDIJINE-8823478; PubMed=2836084; MEDIJINE-8823478; PubMed=2836084; MCCracken A., Uhlenbusch I., Gellissen G "Structure of the cloned Locusta migrator restriction mapping and sequence of its l Curr. Genet. 11:625-630(1987).
EMBL; X05286; CAA28905.1; -.
GO; GO:0005739; C:mitochondrion; IEA.
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P82085;
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01-MAY-2000 (TrEMBLrel 13, I
01-MAY-2000 (TrEMBLrel 13, I
DYNASTIN 7.
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Q34909;
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TISSUB-SKIN SECRETION;
TISSUB-SKIN SECRETION;
Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structure of the dynastins from
"Peptides from Australian frogs. The structure of the dynastins from
Limnodynastes salmini and Fletcherin from Limnodynastes fletcheri.";
Aust. J. Chem. 46:1235-1244(1993).
-!- WASS SPECTROMETRY: MW=1114; METHOD=FAB.
SEQUENCE 12 AA; 1114 MW; 3AB5A976CAA72728 CRC64;
                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
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Bukaryota; Metazoa; Chordata; Cranista; Ve
Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
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-!- MASS SPECTROMETRY: MW=786; METHOD=FAB.
SEQUENCE 8 AA; 786 MW; 7B58772455B05728 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mitochondrion.
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O71066; O71066; O71061
01-AUG-1998 (TremBirel: 07,
01-AUG-1998 (TremBirel: 19,
01-DEC-2001 (TremBirel: 19,
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NON TER
SEQUENCE
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Liermann H., Harder T., Haas L.;
"Genetic analysis of the central un
proximal coding part of the F gene
morbilliviruses.";
Submitted (SEP-1997) to the EMBL/Ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canine distemper virus.
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
NCBI_TaxID=11232;
                                                                                                                                                                                                                                                                                Canine distemper virus.
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
                                                    Liermann H., Harder T., Haas L.;
"Genetic analysis of the central untranslated genome region proximal coding part of the F gene of wild-type and vaccine morbilliviruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; AF028234; AAC00164.1; -.
                                                                                                                                                                         SEQUENCE FROM N.A.
STRAIN=Dog #10757/96;
Liermann H., Harder T
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NCBI_TaxID=11232;
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                                           AF026234;
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ER 9 9
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                     F29D045760440441 CRC64;
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2; Mismatches
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01-NOV-1996 (TrEMBLrel. 01, Cre
01-NOV-1996 (TrEMBLrel. 22, Las
01-OCT-2002 (TrEMBLrel. 22, Las
Alpha-amylase (Fragment).
Oryza satīva (Rice).
Oryza satīva (Rice).
Eukaryota; Viridiplantae; Strep
Spermatophyta; Magnoliophyta; L
Ehrhartoideae; Oryzeae; Oryza.
NCBI TaxID-44530;
                                                                                                                                                      MEDLINE=92013147; PubMed=1717588;
Takahashi K., Kishimoto S., Ohori K., Yoshizawa H., Machida A., Ohonuma H., Tsuda F., Munekata E., Miyakawa Y., Mayumi M.;
"Molecular heterogeneity of e antigen polypeptides in sera from carriers of hepatitis B virus.";
J. Immunol. 147:3156-3160(1991).
PIR; B46453; B46453.
NON TER 10 10
AA; 1238 MW; 485A6E3AB721E9C7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                E antigen P20E (Fragment).

Heparitis B virus.

Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.

NCBI_TaxID=10407;
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01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2003 (TrEMBLrel. 24,
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Rodriquez R.L.
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EMBL; M62916; AAA33892.1;
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yta; Liliopsida; Poales; Poaceae;
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1; Mismatches
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RESULT 13
Q28742
ID Q2874
AC Q2874
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Q85631
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Q23876;
Q1-NOV-1996
Q1-NOV-1996
Q1-NOV-1998
Q28742 PRELIMINARY; PRT; 7 AA. Q28742; Q28742; Q1-NOV-1996 (TrEMBLrel. 01, Created) 01-NOV-1996 (TrEMBLrel. 01, Last sequence update) 01-JUN-2003 (TrEMBLrel. 24, Last annotation update) Alpha-myosin heavy chain (Fragment).
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NON TER
SEQUENCE
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MCKeown M., Firtel R.A.;
"Actin multigene family of Dictyostelium.";
Cold Spring Harb Symp. Quant. Biol. 46:495-505(1982).
EMBL; K02957; AAA33150.1;
EMBL; K02957; AAA33150.1;
EMBL; K02957; AAA33150.1;
SEQUENCE 11 AA; 1205 NW; 728B4C14C6C2CAAB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                        MEDIINE=85033920; PubMed=6092695;
Sutrave P., Jansen H.W., Bister K., Rapp U.R.;
"3'-terminal region of avian carcinoma virus MH2 shares
elements with avian sarcoma viruses Y73 and SR-A.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=11958;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MH2, proviral DNA, myc to 3' LTR (Fragment).
Avian carcinoma virus.
Viruses; Retroid viruses; Retroviridae; Alpharetrovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida;
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K03100; AAA42388.1; -.
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G (TrEMBLrel.
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al DNA, myc to 3' LTR (Fragment).
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O1-MAR-2001 (TrEMBLrel. 16, Cr
O1-MAR-2001 (TrEMBLrel. 24, L;
O1-JUN-2003 (TrEMBLrel. 24, L;
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Q15891;
O1-NOV-1996 (TREMBLrel. 01, 0
01-NOV-1996 (TREMBLrel. 01, I
01-DEC-2001 (TREMBLrel. 19, I
(Clone XP2E8B) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                        "Isolation of chromosome-specific genes arrayed cDNAs and cosmid libraries."; Hum. Mol. Genet. 0:0-0(1995).
EMBL; L32131; AAA73881.1; -.
NON_TER 1 1 1
NON_TER 9 9 9
SEQUENCE 9 AA; 1030 MW; E56635A1A336
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SEQUENCE FROM N.A.
MEDLINE=84221901; PubMed=6328491;
MEDLINE=84221901; Umeda P.K., Sinha
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Eukaryota; Metazoa; (
Mammalia; Eutheria; )
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SEQUENCE 7 AA; 91
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        Linospadix monostachya.
Chloroplast.
Eukaryota, Viridiplantas, Streptophyta, En
Spermatophyta, Magnoliophyta, Liliopsida,
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Lee C.-C., Yazdani A., We
Coolbaugh M.I., Chinault
                                                                             RPS16
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Mammalia; Eutheria;
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ci. U.S.A. 81:3044-3048(1984)
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Primates;
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Lagomorpha; Leporidae; Oryctolagus
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Catarrhini; Hominidae;
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Baldini A., Lindsay F
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OC Areceae; Linospadicinae; Linospadix.

OX NCBI_TaxID=131282;

RN GEQUENCE FROM N.A.

RC TISSUE=Leaf;

RA Asmussen C.B., Chase M.W.;

RT "Coding and noncoding plastid DNA in palm systematics.";

RL Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.

DR GO:0009507; C:chloroplast; IEA.

KW Chloroplast. 1

PT NON_TER 1

FT NON_TER 9

SQ SEQUENCE 9 AA; 1106 MW; 1B9236C2C0441B50 CRC64;

Query Match 28.8%; Score 17; DB 8; Length 9;

Best Local Similarity 75.0%; Pred. No. 1e+06;

Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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Search completed: May 18, 2004, 10:04:47

Job time: 32.25 secs
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Claim 4; Page 92; 119pp; English.

Compounds of the structure or formula S-(L)_n-B, R 1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing CAMP

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

WPI; 2000-452384/39.

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V	Aar91648	7	Aar89301	Abu54029	335	Abg69386	Aay97062	œ	Aab86222	Aar91647	Aab96891	Aab96892	Aap82547	Aab01866	Abp71482	Aaw45785	71	Abp71485	Abp71484
Amino aci	Human par	Fertilin	Japanese	Human DNA	Granulocy	Vascular	PTH-rP N-	PTH(1-14)	Human par	Human par	Rat parat	Rat parat	(Asn10, T	PTH(1-14)	Parathyro	Parathyro	Parathyro	Parathyro	Paratnyro

ALIGNMENTS

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Gardella TJ, Kronenberg HM, Potts JT, Jueppner H;	(JUEP/) JUEPPNER H.	(POTT/) POTTS J T.	(KRON/) KRONENBERG H M.	(GARD/) GARDELLA T J.		31-DEC-1998; 98US-0114577P.	30-DEC-1999; 99WO-US031108.		(06-JUL-2000.)		WO200039278-A2.		Homo sapiens.		bone reformation; resorption; remodeling; tether1; osteoporosis.	PTH; parathyroid hormone; N-terminal; signaling domain; bone mass;		Parathyroid hormone N-terminal signaling domain (residues 1-11).		31-OCT-2000 (first entry)		AAY96968;		AAY96968 standard; peptide; 11 AA.	RESULT 1

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RESULT 2
ABG72607
ID ABG72607
XX ABG7
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XX Huma
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OS Mus
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Best Local 9
The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, being formed from the N-terminus of parathyroid hormone (PTH). Also included are; (1) a method for producing antibodies useful in the determination PTH levels in a biological sample comprising; (a) providing at least on first peptide antigen comprising a peptide fragment of PTH; (b) administering the first peptide antigen to a host animal to induce antibody production; (c) monitoring the antibody titre produced; (d) isolating antisera produced in the host animal; and (e) selecting antisera from the isolated antisera produced in the host that is capable of binding to a second peptide antisera; (2) an antibody (ab) produced be
                                                                                                                                                                                                                                                                                                                                                                                                        New parathyroid hormone (PTH) antigenic peptide inducing the and isolation of antibodies having an affinity to it, useful determining bioactive PTH levels in serum, plasma and/or cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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Sus scrofa
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                                                                                                                                                                                                                                                                                                                                                                                    media.
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y hyperparathyroidism; mouse; rat; bovine; porcine; canine.
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LAVIGNE J R.
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Nismatches
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New

parathyroid hormone (PTH) antigenic peptide inducing the isolation of antibodies having an affinity to it, useful armining bioactive PTH levels in serum, plasma and/or cell

formation for culture

determining

Claim 2;

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English

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                                                                                                                            (ZAHR/)
(LAVI/)
                                                                                                                                                                                                                                                                                                                                              ธนธ
                                                                                                                                                                                                                                                                                                                                                                                Mus
                                                                                                                                                                                                                                                                                                                                                                                                              Human; parathyroid hormone; antigen; hypercalcaemia; osteoporosis; primary hyperparathyroidism; mouse; rat; bovine; porcine; canine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               non-molecular forms of PTH. PTH levels are an important parameter in patients suffering from hypercalcaemia, osteoporosis and primary hyperparathyroidism. The present sequence represents a PTH antigenic peptide representing amino acids 2-12 of human, mouse, rat, porcine,
                                                                                WPI;
                                                                                                      Zahradnik RJ,
                                                                                                                                                                05-DEC-2000; 2000US-00730174
                                                                                                                                                                                      05-DEC-2000;
                                                                                                                                                                                                                                                                       Misc-difference
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                                                                                                                                                                                                              15-AUG-2002
                                                                                                                                                                                                                                     US2002110871-AJ
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                                                                                                                                                                                                                                                                                                                                 scrofa.
s familiaris
                                                                                2003-066685/06
                                                                                                                                                                                                                                                                                                                                                                                          sapiens
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10; Conserv
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LAVIGNE J R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                hormone antigenic peptide 1-12.
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                                                                                                      Lavigne
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Pred. No. 0.012;
0; Mismatches
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The invention relates to a new antigenic peptide for inducing the CC formation and isolation of antibodies having an affinity to it, being CC are; (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one first peptide antigen comprising a peptide fragment of PTH; (b) antibody production; (c) monitoring the antibody titre produced; (d) colating antisera produced in the host animal to induce cc antibody production; (c) monitoring the antibody titre produced; (d) colating antisera produced in the host animal to induce cc antisera from the isolated antisera produced in the host that is capable complicating and (3) test kits and analytical procedures used for the compositions of the present invention are useful for determining cc disolative intact PTH levels in serum, plasma and/or cell culture media. The antigens, antibodies and methods of the present invention, as cc compared to prior art, have the particular advantages of possessing creatives unferring from hypercalcaemia, oscepoporosis and primary completents suffering from hypercalcaemia, oscepoporosis and primary completents suffering from hypercalcaemia, oscepoporosis and primary complete and movine PTH animal and bovine PTH.

8

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VSEIQLMHNLG 12

Query Match Best Local Matches

10;

Conservative

Similarity

84.7%;

Score 50; DB 6; Pred. No. 0.014;

Length 12,

Mismatches

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Sequence 12 AA;

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RESULT 4
AAR91644
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                        New antigenic peptide(s) from human parathyroid hormone - and antibodies generated using them, able to distinguish between active and inactive forms of the hormone.
                                                                                                                                                             DE4434551-A1
                                                                                                                                                 04-APR-1996
                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                    diagnosis;
                                                                                                                                                                                                             Human parathyroid
                                                                                                                                                                                                                              Human parathyroid hormone antigenic peptide hPTH 1-10.
                                                                                                                                                                                                                                                06-NOV-1996
                                                                                                                                                                                                                                                                  AAR91644;
                                                                                                                                                                                                                                                                                  AAR91644 standard; peptide; 10
                                                                            Adermann
                                                                                                               28-SEP-1994;
                                                                                                                                28-SEP-1994;
                                                                                              (FORS/) FORSSMANN W.
                                                            1996-180391/19
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        Page 4; 5pp;
                                                                                                                                                                                                     active hPTH 1-37.
                                                                            Forssmann
                                                                                                                                                                                                                                              (first entry)
                                                                                                              94DE-04434551.
                                                                                                                                94DE-04434551
                                                                                                                                                                                                            hormone; hPTH; antigen; alpha-helix; immunogen;
                                                                            Hock
                                                                            ā
                                                                            Maegerlein
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B 5

Matches Query Match Best Local

10;

Conservative

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Mismatches

Indels

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Gaps

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Similarity

83.1%;

Score 49; ; Pred. No.

DB 3; 0.017;

Length 10;

1 SVSEIQLMHN 10

SVSEIQLMHN

Sequence 10 firming

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RESULT 5
AAY68767
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Best Local
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                                    The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipophilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin
                                                                                                                                                                                              New parathyroid hormone fragment peptides, used as lipolysis stimulants in topically applied cosmetic compositions for slimming treatment of excessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                                                  27-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY68767;
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                                                                                                                                                                                                                                                                                                                                                                      07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amino acids 1-10 of a parathyroid hormone (pTH).
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                                                                                                                                                                  Claim 1; Page 8; 18pp; French.
                                                                                                                                                                                                                                                        WPI; 2000-171243/15.
                                                                                                                                                                                                                                                                                                                                           17-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                           WO200004047-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    slimming treatment; cellulite; skin firming.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parathyroid hormone; lipolysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-MAY-2000
                                                                                                                                                                                                                                                                                   Lintner K;
                                                                                                                                                                                                                                                                                                                (SEDE-) SEDERMA
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Pred. No.
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0.017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dermatological; skin care;
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RESULT 6
AAB86219
ID AAB862219
ID O3-S
XX AAB8
AX AAB8
AX O3-S
XX Para
AX Para
RESULT 7
ABR44166
ID ABR4
XX
AC ABR4
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calciun-metabolism disorders, osteopathy, or hypo- or hyper-parathyroidism. The method (unlike known assays) recognizes that (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically cinactive, and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hyper-parathyroidism.
                                 ABR44166
                                                                                           ABR44166 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-376318/40
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                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
nes 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
                                                                                                                                                                                                                                                                                                                   SVSEIQLMHN 10
                                                                                                                                                                                                                                                          SVSEIQLMHN 10
                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                   83.1%; 5c.
100.0%; Pr
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. 0.017;
thes 0;
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AABB62
ID AABB
XX AABB
XX AABB
XX AABB
XX PAIR
CT 03-S
XX PAIR
CK PAIR
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KW DELD
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PF N DELD
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Best Local
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Chang M;
                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a fusion peptide (Tat-hPTHDP), where self cell-penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmetic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endothelium, does not cause skin disease and has superior lipolysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
                                                                                         Homo
                                                                                                                             Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                                                                                                   Human parathyroid
                                                                                                                                                                                              03-SEP-2001
                                                                                                                                                                                                                                              AAB86225 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel fusion peptide comprising self to human parathyroid hormone-derived slimming cosmetic composition.
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15-MAR-2002; 2002KR-00014062.
                                                                                                                hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                                                                                      AAB86225
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                   hormone
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                                                                                                                                                                                                                                                                                                                                     10
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                                                                                                                                                                                                                                                                                                                                                           J.0%;
                                                                                                                                                                   immunogenic peptide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cho W,
                                                                                                                                                                                                                                                                                                                                                               Score 49; DB; Pred. No. 0.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fragment (residues 1-10).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell-penetrating Tat peptide bound peptide, useful as component of skin
                                                                                                                                                                                                                                                                                                                                                           DB 6; be...
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                                                                                                                                                                                                                                                                                                                                                                                        Length 10
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21-JUN-2001.

DE19961350-A1

17-DEC-1999;

99DE-01061350

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RESULT 9
AAR91647
ID AAR9
XX AAR9
AC AAR9
AC AAR9
XX Huma
XX Huma
XX Huma
XX Huma
XX Synt
XX DE44
XX DE44
XX DE44
XX E8-S
XX 28-S
XX 28-S
XX Y Ade1
XX WPI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC This invention describes a novel method for determining (M1) the content CC of active parathyroid hormone (A) by treating a sample with (i) antibody CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to Calculate the content of physiologically active (A). The method is used for diagnosis of calcium-methodism disorders, osteopathy, or hypo- or CC hyper-parathyroidism. The method (unlike known assays) recognizes that CC (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically CC active and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This conserver, and the content of physiologically active (A); contrast methods to illustrate the method of the inverse of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This
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Matches
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                                                                                                                          28-SEP-1994;
                                                                                                                                                              28-SEP-1994;
                                                                                                                                                                                                04-APR-1996
                                                                                                                                                                                                                                        DE4434551-A1
                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                 Human parathyroid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-376318/40
                                                                                    (FORS/) FORSSMANN W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention
                                                                                                                                                                                                                                                                                                                                                                         parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          h 76.3%;
Similarity 100.0%;
9; Conservative
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                                                                                                                        94DE-04434551
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                                                                                                                                                                                                                                                                                                                                   hormone; hPTH; antigen; alpha-helix; immunogen
                                                                                                                                                                                                                                                                                                                                                                       hormone antigenic peptide hPTH 1-9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 9;
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WPI; 1996-180391/19

Adermann K,

Forssmann

Σ,

Hock D,

Maegerlein M;

Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having FTH-1 receptors. Or screening for a peptide or

or S-(L)_n

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

WPI; 2000-452384/39

Claim 11; Page 93; 119pp; English.

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AAY96981
ID AAY96
XX AAY96
XX AAY96
XX DT 31-OC
XX PTH;
XW DONE
XX WO200
XX ICAND
PA (FOTTI
XX WPT;
XX WPT;
XX New PT Linke
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GARD/) GARDELLA T (KRON/) KRONENBERG F (POTT/) POTTS J T. (JUEP/) JUEPPNER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a specific example of claimed immunogenic peptides having a sequence from hpTH(1-37) which includes the N- or C-terminal alpha-helical region and/or the non-structured region of the hormone. Antibodies and their binding fragments generated by injecting animal with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tether1; osteoporosis.
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Pred. No. 1.4e+06;
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(claimed). The new compound can be administered ke the large native PTH or PTHrP which avoids the ons to treat osteoporosis

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                                                          This invention describes a novel method for determining (M1) the content cof active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that recognizes an epitope within the receptor-binding site of (A). The number color of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or hyper-parathyroidism. The method (unlike known assays) recognizes that c(i) some fragments of (A) shorter than the complete (4) as a peptide are active and (ii) that apparently intact peptide may be biologically inactive, and also takes into account the fact that some fragments of (A) contrast methods that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
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Pred. No. 1.4e+06;
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This invention describes a novel method for the preparation of peptides (CI) that contain both cyclic and linear peptide fragments comprises sequential reaction of a resin-bound linear fragment with the Cyclic CC fragment in N-protected form and optionally other linear fragments. The products of the invention have osteopathic and hypotensive activity. (II) CC bind to hpTH receptors and act as agonists or antagonists of hpTH. The CC method is particularly used to prepare cyclic peptide analogs of CC parathyroid hormone (PTH) or PTH-related peptides which are useful for CC treating diseases that respond to treatment with agents that bind to PTH ceceptors (with or without activation of adenylyl cyclase activity), e.g. hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, cushing's syndrome, renal failure and hypertension, also for promoting repair of bone fractures. Separate synthesis of the cyclic fragment allows convergent synthesis of resin-bound (II), with better cyilds and higher throughput. The difficulties associated with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 75; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Convergent synthesis of peptides for treating e.g. bone disorders
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The invention relates to a novel parathyroid hormone (PTH) peptide (AAB01853) and parathyroid hormone-related peptide (PTHP), AAB01861), and biologically active derivatives thereof (AAB01857-B01858, AAB01861).

B01869). The peptides of the invention are at least 8% identical to the generic peptide of the formula: X1-Val-Ser-Glu-X2-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is 1le or Lys; X3 is Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; provided that the peptide is not PTHP(1-14). The peptides of the invention consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-terminal derivatives. PTH is a major regulator of calcium homeostasis, and is necessary for the normal function of the gastrointestinal, skeletal, neurological system, neuromuscular and cardiovascular systems. It binds to both PTH-1 receptors on osteoblasts and renal tubular cells, and to the recently identified PTH-2 receptor. PTH has a potent anabolic effect on the skeleton, and mediates calcium reabsorption, enhances of the renal and skeletal actions of PTH, and also bind to the PTH-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Matches 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 26; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Parathyroid hormone (PTH) peptides, PTH related peptides and the nucleic acids that encode them, useful for treating osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GARD/)
(KRON/)
(POTT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-339693/29
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KRONENBERG H
POTTS J T.
JUEPPNER H.
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83.3%;
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                                                     receptor. They do not bind to the PTH-2 receptor. The peptides of the invention are either agonists of pTH-1 and pTH-2 receptors (AAB01858, AAB01861-B01866) or are PTH-1/PTH-2 receptor antagonists (AAB01867-B01869). PTH-1/PTH-2 receptor agonists are useful for the treatment of conditions characterised by a decrease in bone mass, such as osteoporosis. PTH-1/PTH-2 receptor antagonists are useful for treatment of periodical disorders that arise from excessive or altered action of the PTH-1/PTH-2 receptor. Detectably labelled peptides of the invention are also useful in the determination of rates of bone formation, bone resorption and/or bone remodelling in a patient. The peptides of the invention are "minimised" versions of PTH or PTH-P which are inexpensive to prepare by conventional synthetic chemistry, and can be delivered to a patient via non-parenteral routes. Sequences AAB01858 and AAB01861-B01866 represent PTH-1/PTH-2 receptor agonists
Sequence 9
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Matches
                                        Query Match
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8; Conserv
AVSEIQLMH
            SVSEIQLMH
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88.9%;
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Pred. No. 1.4e
1; Mismatches
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1.4e+06;
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RESULT 14
AAY96966
PTH;
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                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                          AAY96966 standard;
                                                                                                                                                                              Parathyroid hormone N-terminal signaling domain (residues 1-9).
                                                                                                                                                                     parathyroid hormone; N-terminal; signaling
                                                                                                                                                                reformation;
                                                                                                                                                                                       (first entry)
                                                                                                                                                                resorption;
                                                                                                                                                                                                         peptide;
                                                                                                                                                                remodeling; tether1;
                                                                                                                                                                     domain;
                                                                                                                                                                      bone mass;
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30-DEC-1999; 06-JUL-2000 99WO-US031108

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(KRON/)
 (JUEP/)
                            (GARD/)
) KRONENBERG H M.
) POTTS J T.
) JUEPPNER H.
                            GARDELLA
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31-DEC-1998;

98US-0114577P

Gardella TJ, Kronenberg , MH Potts JT, Jueppner

2000-452384/39

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass. Claim 4; Page 92; 119pp; English

Compounds of the structure or formula S-(L) n-B, R-1-S-(L) n-R or S-(L) n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a postitive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R-1 is the PTH-1 receptor signal sequence; and R is (a portion of PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases

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The invention relates to a novel parathyroid hormone (PTH) peptide (CAAB01859) and parathyroid hormone related peptide (PTHFP, AAB01860), and biologically active derivatives thereof (AAB01857-B01858, AAB01861). The peptides of the invention are at least 83 identical to the (C generic peptide of the formula: X1-Val-Ser-GlN-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is lle or Lys; X3 is CC Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; Cl provided that the peptide is not PTHFP1-14). The peptides of the cornisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-C consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-C cand is necessary for the normal function of the gastrointestinal, CC skeletal, neurological system, neuromuscular and cardiovascular systems. CC it binds to both PTH-1 receptors on oscoblasts and renal tubular cells, and the recently identified PTH-2 receptor. PTH has a potent anabolic effect on the skeleton, and mediates calcium reabsorption, enhances companies to the clearance and vitamin D synthesis in the kidney. A homologous
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthesis; bone synthesis; agonist; osteoporosis; non-parenteral delivery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and disorders associated with decreased tether1 activity, increas in a mammalian cell having PTH-1 receptors, or screening for a per non-peptide PTH (claimed). The new compound can be administered be inhalation unlike the large native PTH or PTHrP which avoids the
                                                                                                                                                                                                                                                                                                                   Disclosure; Page 26;
                                                                                                                                                                                                                                                                                                                                                 Parathyroid hormone (PTH) peptides, PTH related peptides and the nucleic acids that encode them, useful for treating osteoporosis.
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(KRON/)
(POTT/)
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88.9%;
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Pred. No. 1.4e+06;
1; Mismatches 0
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Search completed: May 18, 2004, 10:01:53 Job time : 46.75 secs

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                                                                                                                                                                                                                                                                                                                                                                              calls of protein hormones, the PTH-related proteins (PTHrP) mimic some of continuous control of the renal and skelatal actions of PTH, and also bind to the PTH-1 control of the PTH-1 control of the protein of the protein of the protein of the control of the control of the protein of the pro
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AVSEIQLLH
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Result
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Maximum
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Maximum
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/ Cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
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                 US-09-730-174A-5
US-09-730-174A-1
US-09-730-174A-1
US-09-730-174A-6
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US-09-730-174A-6
US-09-730-174A-2
US-10-168-185-7
US-10-168-185-7
US-10-168-185-7
US-10-168-185-7
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Sequence 5, Appli
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US-09-730-174A-5
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Patent No. US20020110871A1
GENERAL INFORMATION;
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT PILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local S
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equence 248, Ap	quence 83, App	equence 31,	equence 49, App	псе 288, Ар	equence 249	e 50,	Ð	equence 11,	ce 4,	quence 332,	e 12,	equence 12,	nce 469	equence 294	quence 41,	equence 41,	quence 41,	equence 41,	equence 41,	equence 41,	equence 369	quence 27,	equence 185, Ap	equence 11, App	equence 5,	equence 16, App	6, A	equence 61, App	

ALIGNMENTS

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Sequence 5, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Pel
TITLE OF INVENTION: Selective Bindin;
FILE REFERENCE: IMUNE-001A
                        and Peptide Antigens for Producing Antibodies having Binding Specificity to Bioactive Intact Parathyroid
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Pred. No. 0.00071;
; Mismatches 0;
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of Antibodies and Isolates

said

Antibodies

Length 12

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RESULT 4
US-09-730-174A-4
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US-09-730-174A-1
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                                                                                                                   TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H FILE REFERENCE: IMUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12
SEQ ID NO 4
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zahradnik, R.J.

TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid HC FILE REFERENCE: IMUNE-001A CURRENT FILING NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12

SEQ ID NO 1

LENGTH: 11

TYPE: PRT
ORGANISM: Artificial Sequence
Query Match 93.2%;
Best Local Similarity 91.7%;
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
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Best Local (
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 5
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Best Local Similarity
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                            OTHER INFORMATION:
                                                                                                                  FEATURE:
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1 VSEIQLMHNLG 11
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100.0%;
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Score 55; DB 9; Li
Pred. No. 0.0036;
0; Mismatches 1;
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Pred. No. 0.003
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Pred. No. 0.0024;
1; Mismatches
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                                                                                            of Antibodies and Isolates
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                                                                  Sequence 1, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
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RESULT 7
US-10-168-185-1
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US-09-730-174A-2
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US-09-730-174A-6
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Patent No. US20020110871A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                  SEQ ID NO 2
LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hopert Reference: IMUNB-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
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Best Local S
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Best Local Similarity 83.
Matches 10; Conservative
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Patent No. US20020110871A1
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Antibodies and Peptide Antigens fo
TITLE OF INVENTION: Selective Binding Specificity to
FILE REFERENCE: IMUNE-001A
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Zahradnik, R.J
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TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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Pred. No.
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Pred. No. 0.012;
1; Mismatches
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APPLICANT: Armbruster, Franz Paul APPLICANT: Missbichler, Albert APPLICANT: Schmidt-Gayk, Heinric APPLICANT: Roth, Heinz-Jurgen

Schmidt-Gayk, Heinrich

Heinz-Jurgen N: Method for Determining

Parathormone

TITLE OF INVENTION:

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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Method for Determining Parathormone
FILE REFERENCE: HLZ-004US
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 2001-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FITLE OF INVENTION: Activity in a Human Sample FIL3 REFERENCE: HLZ-004US
CURRENT APPLICATION UNMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: ECT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
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US-10-168-185-7
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Best Local S
Matches 9
                                                                                                                                                                                          Sequence 2, Application US/10168185 Publication No. US20030175802A1 GENERAL INFORMATION:
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Best Local :
                                         APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Albert
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 9
TYPE: PRT
ORGANISM: Homo sapiens
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TYPE: PRT
ORGANISM: Homo sapiens
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les 9; Conserv
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100.0%; Pr
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APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REPERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 1909-12-17
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SEQ ID NO 2
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Publication No. US20030166838A1
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
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Publication No. US20030175802A1
GENERAL INFORMATION:
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Best Local 9
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Matches 9; Conserv
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PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives
TITLE OF INVENTION: Broactive Peptides (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH)
FILE REFERENCE: 0609.4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
                                                                                                                                                                      APPLICANT: Gardella, Thomas J. APPLICANT: Kronenberg, Henry APPLICANT: Potts, John T. APPLICANT: Juppner, Harald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Armbruster, Franz Paul APPLICANT: Missbichler, Albert
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TYPE: PRT
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Schmidt-Gayk, Heinrich
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                                                                                                                                                                                                                                                 a, Thomas J.
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Sequence 7, Application US/10192673

Publication No. US20030166638A1

GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: With Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)

FILE REFERENCE: 0609.4570002

CURRENT APPLICATION NUMBER: US/10/192.673

CURRENT APPLICATION NUMBER: US. 09/421.379

FRIOR APPLICATION NUMBER: U.S. 09/421.379

PRIOR APPLICATION NUMBER: U.S. 09/105.530

PRIOR APPLICATION NUMBER: U.S. 60/105.530

PRIOR APPLICATION SERVICE 
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US-10-192-673-7
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Best Local Similarity
Matches 8; Conserv
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                                                        APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidd-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1999-10-20 PRIOR APPLICATION NUMBER: U.S. PRIOR FILING DATE: 1998-10-22 NUMBER OF SEQ ID NOS: 13
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 77.
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88.9%;
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77.8%;
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2; Mismatches
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Pred. No. 1e+06;
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APPLICANT: Kronenberg, Henry
APPLICANT: Kronenberg, Henry
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Bioactive Peptides and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT APPLICATION NUMBER: US/0/192,673
CURRENT FILING DATE: 2002-07-11
FRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR PILING DATE: 1999-10-20
PRIOR PILING DATE: 1999-10-20
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
LENGEL: 0
      ; Sequence 4, Application US/10168185
; Publication No. US20030175802A1
; GENERAL INFORMATION:
   APPLICANT: Armbruster, Franz Paul
   APPLICANT: Missbichler, Albert
   APPLICANT: Schmidt-Gayk, Heinrich
   APPLICANT: Roth, Heinz-Jurgen
; ATTLE OF INVENTION: Method for Determining Parathormone
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Best Local Similarity
7; Conserve
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CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-168-185-4
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TYPE: PRT
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TYPE PRINCIPLE PRINCIPLE SEQUENCE PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
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8; Conserv
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77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 33; DB 14; Length 9 Pred. No. 1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT APPLICATION NUMBER: US/2-06-17
PRIOR APPLICATION NUMBER: DCT/EP00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILLING DATE: 199912-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 7
TYPE: PRT
ORGANISM: Homo sapiens
US-10-168-185-4

Query Match
Best Local Similarity 100.0%; Pred. No. 1e-06;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0,

Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps O;

Search completed: May 18, 2004, 10:20:59

Job time: 33.25 secs
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Result
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Maximum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OM protein - protein search,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         on:
           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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DB
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           Query
Match
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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    US-09-442-989-31

US-08-817-547A-1

US-08-817-547A-1

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US-09-421-379-6

US-09-421-379-10

5460978-3

US-08-817-547A-4

US-08-817-547A-14

US-08-817-547A-14

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-706-56

US-08-817-547A-15

US-08-817-547A-15

US-08-817-547A-13

US-08-817-547A-13

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US-08-817-547A-13

US-08-817-547A-13

US-08-817-547A-13

US-08-817-547A-13

US-08-817-547A-13

US-08-318-693-13

US-08-318-082B-41

US-09-336-093-12
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Sequence 31, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3, Appli Sequence 3, Appli Sequence 10, Appli Sequence 4, Appli Sequence 4, Appli Sequence 15, Appli Sequence 15, Appli Sequence 17, Appli Sequence 16, Appli Sequence 17, Appli Sequence 17, Appli Sequence 18, Appli Sequence 11, Appli Sequence 11, Appli Sequence 14, Appli Sequence 14, Appli Sequence 13, Appli Sequence 14, Appli Sequence 13, Appli Sequence 14, Appli Sequence 13, Appli Sequence 14, Appli Sequence 14, Appli Sequence 12, Appli Sequence 14, Appli Sequence 14, Appli Sequence 15, Appli 
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US-09-442-989-31
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ALIGNMENTS

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US-08-817-547A-1
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Sequence 2, Application US/08817547A
PATENT NO. 6030790
GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application Patent No. 6030790
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Best Local
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Best Local
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TELEPHONE: 404-818-3790
TELEPAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: PCT/EP95/
APPLICATION NUMBER: PCT/EP95/
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: unknownouscure Type; po
HYPOTHETICAL: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acida STAANDENNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/817,547A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                              1 SVSEIQLMHN 10
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10; Conserv
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100.0%; Pred. No. 0.0057;
tive 0; Mismatches 0
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Pred. No. 0.0045;
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                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Konenberg, Henry
APPLICANT: Konenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
ITITE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
ITITE OF INVENTION: Bloactive Peptides (PTH) and Parathyroid
ITITE OF INVENTION: Hormone-Related Peptide (PTHrp)
FILE REFERENCE: 0609,4570001
FILE REFERENCE: 0609,4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER APPLICATION NUMBER: US. 50/105,530
EARLIER APPLICATION NUMBER: US. 50/105,530
EARLIER APPLICATION NUMBER: US. 50/105,530
SEGID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEG ID NOS: 13
    ; ORGANISM: Artificial Sequence; FEATURE:
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-09-421-379-6
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 22,176
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 6, Application US/09421379
Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO.
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                     TYPE: PRI
                                                                                                                      ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: unknown
MOLECULE TYPE: po
HYPOTHETICAL: no
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MEDIUM TYPE: Floppy disk
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TYPE: amino acid
STRANDEDNESS: unknown
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ZIP: 30303
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SVSEIQLMH 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72.9%; Score 43; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
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3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 9;
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Gaps

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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Notes, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0509.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CARLIER APPLICATION NUMBER: US/09/421,379
EARLIER APPLICATION NUMBER: US/09/421,379
EARLIER FILING DATE: 1999-10-20
EARLIER FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTMARE: Patentin Ver. 2.0
SEQ ID NO 7
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      뭥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-817-547A-3
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                                                                                                                                                                                                                                                                             Hatent NO. BURNERS INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
APPLICANT: Magerlein, Markus
APPLICANT: Magerlein, Peptides from the hPTH Sequence
TITLE OF SERUJENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 88.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA
                                                                                                                                                                                                                                 STREET: 191 PORTION OF CITY: Atlanta
                                 APPLICATION NUMBER: US/08/817,547A
                                                                                                                                                                         ZIP: 30303
                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 7; Conserv
                                                                                                                                                                                              Georgia
Y: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AVSEIQLLH 9
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                                                                                                                                                                                                                                                  E: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
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77.8%;
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Pred. No. 3e+0
2; Mismatches
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Pred. No. 3e+05;
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Fotts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Broactive Peptides and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTHP) and Parathyroid
TITLE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOPTWARE: Patentin Ver. 2.0
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                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-421-379-10
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US-08-817-547A-3
                                                                 Query Match
Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                         LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 078
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: ur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: unl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/E
FILING DATE: 29 SEPT 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10,
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                             1 SVSEIQLMH 9
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                                                                 Similarity 77.4
7; Conservative
SVSEHQLLH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09421379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 amino acids
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Conservative (
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                                                                                  55.9%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      59.3%; Score 35; DB 1100.0%; Pred. No. 3e-
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                                                                 Score 33; DB 4;
Pred. No. 3e+05;
1; Mismatches
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                                                                                                  Length 9;
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                                                               Gaps
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5460978-3
; PATENT NO. 5460978
; PATENTI MARTIN, THOMAS J.; MOSELEY, JANE M.; KEME,
; BRUCE E.; WETTENHALL, RICHARD E.H.
; TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL
; HYPERCALCEMIA OF MALIGNANCY-PTHRP

RESULT 8

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RESULT 9
US-08-817-547A-4
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                                                                                                  ; HYPOTHETICAL:
; ANTI-SENSE: 1
US-08-817-547A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/08817547A Patent No. 6030790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 54.2%;
Best Local Similarity 70.0%;
Matches 7; Conservative
                                                 Query Match
Best Local :
                                Matches
                                                                                                                                                                                                                                                                                  PILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: PCT/EP95/03757
RILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-18-3700
                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Adermann, Knuc
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 199,235
FILING DATE: 09-MAY-1988
                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DABER: US/07/715,280
APPLICATION NUMBER: US/07/715,280
FILING DATE: 14-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: unk
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
1 SVSEIQL
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                              Similarity 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
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                             Conservative
                                                                                                                    no
                                                                                                                                  no
                                                                                                                                                   peptide
                                                                                                                                                                                   unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptides from the hPTH Sequence
                                               50.8%; Score 30; DB 3; Length 7; 100.0%; Pred. No. 3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/817,547A
                              0;
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Pred. No. 8.2;
1; Mismatches
                                Mismatches
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                                0; Indels
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                              Gaps
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GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Cardella, Thomas J.
APPLICANT: Notes, John T.
APPLICANT: Notes, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Hormone-Related Peptide (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
CURRENT SPILICATION UNUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER FILING DATE: 1999-10-20
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN DATE: 198-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 8
LENGTH: 9
TYPE: PRT
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                                                                                   Matches
                                                                                                                      Query Match
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Patent No. 6495662
                                                                                                                                                                                   FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 μ
1 AVSEHOLLH 9
                                  1 SVSEIQLMH 9
                                                                                 Similarity 66.7
6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SVSEIQL
                                                                                                    50.8%;
                                                                                                    Score 30; DB 4;
Pred. No. 3e+05;
                                                                                   Mismatches
                                                                                                                           DB 4; Length 9,
                                                                                   Indels
                                                                                   0,
                                                                                   Gaps
                                                                                   0
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WESULT 11

US-08-817-547A-5

(Sequence 5, Application US/08817547A

Patent No. 6030790

PATENT NOCKMATION:

APPLICANT: Adermann, Knut

APPLICANT: Mock, Dieter

APPLICANT: Magerlein, Markus

TITLE OF INVENTION: Peptides from the hPTH Sequence

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Jones & Askew, LLP

STREET: 191 Peachtree Street, 37th Floor

CITY: Atlanta

STATE: Georgia
COUNTRY: USA
ZIP: 30303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 22,176
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007

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COUNTRY: USA
ZIP: 10016-2391
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/428,257A
FILING DATE: 07/05/95
CLASSIFICATION: 514
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14. Application US/08428257A

Patent No. 5885808

GENERAL INFORMATION:
APPLICANT: Sponer, Robert A.
APPLICANT: Epenetos, A.A.
ITILE OF INVENTION: Compounds to target cells
NUMBER OF SEQUENCES: 80
CORRESPONDENCE ADDRESS:
ADDRESSES: Jules 3. Goldberg
STREET: 261 Madison Avenue
CITY: New York
STATE: NY
                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 12
US-08-428-257A-14
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                                                                                                       RESULT 13
US-08-802-981-124
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                                                                                                                                                                                                                                                                                                                                  LENGTH: 10 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-428-257A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-817-547A-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local !
                                                                     Sequence 124, Application US/08802981 Patent No. 6037137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOCE TYPE: per MOLECULE TYPE: per
APPLICANT: Komoriya, Akira
APPLICANT: Packard, Beverly S.
TITLE OF INVENTION: Compositions for the Detection of Enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14.1%; Score 26; DB 3; Local Similarity 100.0%; Pred. No. 3e+05; res 6; Conservative 0: Miematahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 6 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 SVSEIQ 6
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                                                                                                                                                                                                                                                        44.1%; Score 26; DB 2; Length 10; 71.4%; Pred. No. 94; tive 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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APPLICATION NUMBER: US/08/802,981
FILING DATE: 20-FEB-1997
CLASSIFICATION: 435
ATTORNBY/AGENT INFORMATION:
NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 016865-000300US
TELECOMMUNICATION: INFORMATION:
TELEPAN: (415) 576-0200
TELEFAX: (415) 576-0200
TELEFAX: (415) 576-030
INFORMATION FOR SEQ ID NO: 124:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                               FEATURE: Modified-site
NAME/KEY: Modified-site
LOCATION: 3
LOCATION: 3
OTHER INFORMATION: /product= "Aib"
US-08-802-981-124
                                            Matches
                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: #10ppy disk
COMPUTER: IBM PC compatible
COMPUTER: FORM:
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: PATENTIA PC-BOSS
SOFTWARE: PATENTIA PC-BOSS
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                              Match 42.4%; Score 25; DB 3; Local Similarity 80.0%; Pred. No. 1.6e+02; es 4; Conservative 1; Mismatches o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Francisco
                                                                                                                                                                                                                                                                                        STRANDEDNESS:
6 QLMHN 10
                                                                                                                                                                                                                                                                                                           amino acid
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                                                                                       Length 11;
                                            Indels
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RESULT 14
US-08-177-109A-56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 56, Application US/08177109A; Patent No. 5869615; GENERAL INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/177,109A
FILING DATE: 03-JAN-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dennis E. Hourcade and Teresa J. Oglesby TITLE OF INVENTION: MODIFIED COMPLEMENT PROTEASES NUMBER OF SEQUENCES: 62
                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                  ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
CITY: Atlanta
STATE: Georgia
                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                       30309-3450
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RESULT 15
US-08-687-706-56
; Sequence 56, Application US/08687706
; Patent No. 5928992
; GENERAL INFORMATION:
    APPLICANT: Dennis E. Hourcade and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ş
                                                                                       US-08-687-706-56
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                                                                                                                                                                                    APPLICATION NUMBER: US/08/687,706
FILING DATE: 26-UUL-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/177,109
FILING DATE: 03-UAN-1994
CLASSIFICATION: 514
ACCLASSIFICATION: 514
ACCLASSIFICATION: 514
ACCLASSIFICATION: 10FORMATION:
NAME: Pabst, Patriea I.
NAME: Pabst, Patriea I.
REGISTRATION NUMBER: WU 107 DIV
REFERENCE/DOCKET NUMBER: WU 107 DIV
TELEPHONE: (404) 873-8794
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACCTERISTICS:
LENGTH: 5 aning acids
TYPE: acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-177-109A-56
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REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: WU 107
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
TELEPHAX: (404) 873-8795
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
Query Match 40.7%; Score 24; DB 2; Length 5; Best Local Similarity 60.0%; Pred. No. 3e+05; Matches 3; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 40.7%;
Best Local Similarity 60.0%;
Matches 3; Conservative
                                                                                                       TOPOLOGY: 1i:
MOLECULE TYPE:
HYPOTHETICAL: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 30309-3450
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Dennis E. Hourcade and Teresa J. Oglesby
TITLE OF INVENTION: MODIFIED COMPLEMENT PROTEASES
NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
ADDRESSE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
CITY: Atlanta
STATE: Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                          LENGTH:
TYPE: a
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                                                                                                                                               amino acid
                                                                                                       NO NO
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0; Indels
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Gaps
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Qy 8 MHNLG 12
| | | : | 12
| Db 1 LHNMG 5
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Search completed: May 18, 2004, 10:06:43 Job time: 12.75 secs

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Result
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Maximum Match
Listing first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OM protein - protein search, using sw model
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            seq
                                      of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          length: 0 length: 12
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              May 18, 2004, 09:56:10 ; Search time 10.25 Seconds (without alignments) 112.614 Million cell updates/sec
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61
1 SVSEIQFMHNLG 12
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                                    Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                        1111 0111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00111 00
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t 45 summaries
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S35490
PH1190
S51737
A35779
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A32428
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                                                                                                                                                                                                                         amine oxidase (cop
polyphosphate-gluc
Ig heavy chain CDR
N-methylpurine DNA
alpha-2-macroglobu
                                                                                                           rhodopsin homolog
type II site-speci
T-cell receptor al
T-cell receptor be
                                                                                                                                                                                                                                                                                                                                                                                                                                                           carnocin UI49 - Ca
ribosomal protein
sperm-activating p
neuropeptide M-I
                                                    neuropeptide Antho
peptidyl-dipeptida
ribosomal protein
                                                                                                                                                                                                                                                                                                                         hypertrehalosemic hypertrehalosemic
                                                                                                                                                                                                                                                                                                                                                                                                  Ig heavy chain C r
Ig heavy chain V r
T-cell receptor al
adipokinetic hormo
neuropeptide Led-C
                  hypertrehalosemic adipokinetic horm
                                                                                                                                                                                      hypothetical prote
lectin GNL1 alpha
                                                                                                                                                                                                                                                                                                                                                               alpha-myosin heavy
kidney and bladder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytochrome b559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Gene: psbF
A;Genome: chloroplast
C;Superfamily: cytochrome b559 component
C;Keywords: chloroplast; photosynthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytochrome b559 component psbF - pepper chloroplast (fragment)
C;Species: chloroplast Capsicum annuum (pepper)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 11-Jun-1999
C;Accession: S28055
R;Kuntz, M.; Camara, B.; Weil, J.H.; Schantz, R.
Plant Mol. Biol. 20, 1185-1188, 1992
A;Title: The psbL gene from bell pepper (Capsicum annuum): plastid RNA editing also A;Reference number: S28055; MUID:93099270; PMID:1463853
A;Accession: S28055
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S28055
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-10 < KUN>
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Best Local S
Matches 4
                                                                           Matches
                                                                                                             Query Match
                                                                                           Local
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1 SISAMQFI 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4;
                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
SEIQ 5
                                  SEIQ 6
                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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F photosystem II; thylakoid

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DB 2;

Length 10;

ALIGNMENTS

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C;Accession: A58718
R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, I.F.
Appl. Environ. Microbiol. 58, 1417-1422, 1992
A;Title: Purification and characterization of a new bacteriocin isolated from a Carnoback R;Reference number: A58718; MUID:92321768; PMID:1622206
A;Accession: A58718
                                                                          A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 <5TO>
C;Keywords: antibiotic; lanthionine
                                                                                                                                                                                                                                                                                                     RESULT 2
A58718
carnocin UI49 - Carnobacterium sp. (fragment)
                                                                                                                                                                                                                                                             C;Species: Carnobacterium sp.
C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
100.0%; 5
                      29.5%; Score 18;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 23; DB
Pred. No. 2.1e
3; Mismatches
      Mismatches
DB 2; ____, 2.8e+05; 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1e+02;
                                        Length 7;
      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
      0
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      Gaps
      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurit Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and otus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Reference number: A60527
                                                                                                                                      A;Residues:
C;Keywords:
                                                                                                                                                                                                R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rinehart Jr., Biochem. Biophys. Res. Commun. 124, 350-358, 1984
A;Title: Structures of two cockroach neuropeptides assigned by fast atom bombardment A;Reference number: A90118; MUID:85046530; PMID:6548628
A;Accession: A05169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: S78420
R;Goldschmidt-Reisin, S.; Graack, H.R.
submitted to the Protein Sequence Database, February 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ribosomal protein RL41, mitochondrial [validated] - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000
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S78420
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C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #t
C;Accession: A05169
                                                                                                                                                                                                                                                                                                                                          neuropeptide M-I - American cockroach
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C;Species: Heterocentrotus mamillatus
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
C;Accession: A60589
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A;Residues: 1-9 <GOL>
A;Note: the protein is designated as mitochondrial ribosomal protein
C;Keywords: mitochondrion; protein biosynthesis; ribosome
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A;Accession: S78420
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1; Mismatches
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Pred. No. 2e+03;
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A;Molecule type: mRNA
A;Residues: 1-10 <RUF:
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A; Residues: 1-12
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T-cell receptor alpha chain V region (Cw3/1F11) - human (fragment)
(;Species: Homo sapiens (man)
(;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
(;Accession: PH1187
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra,
J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell
A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1187
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R;Makiya, R.; Stigbrand, T.
Eur. J. Biochem. 205, 341-345, 1992
A;Title: Placental alkaline phosphatase has a binding site A;Reference number: S21205; MUID:92209522; PMID:1555592
A;Accession: S21205
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221205

19 heavy chain V region - human

C;Species: Homo sapiens (man)

C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995

C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995
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C;Species: Mus musculus (house mouse)
C;Date: 19-Mar:1997 #sequence_revision 13-Mar-1998 #text_change
C;Accession: A37268
R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R.
J. Biol. Chem. 266, 6607-6613, 1991
A;Tile: Heavy and light chain variable region sequences and ant
A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: A37268
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C;Keywords: heterotetramer; immunoglobulin
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1; Mismatches 4
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Pred. No. 3.8e+03;
1; Mismatches
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A; Residues: 1-10 <GABL-,
A; Residues: 1-10 <GABL-,
A; Residues: 1-10 <GABL-,
R; Gaede, G.; Rinehart Jr., K.L.
R; Gaede, G.; Rinehart Jr., K.L.
Biol. Chem. Hoppe-Seyler 368, 67-75, 1987
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 05-Nov-1999
C;Accession: 146868
R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H.
Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984
A;Title: Characterization of genomic clones specifying rabbit alpha- and bet A;Reference number: 146868; MUID:84221901; PMID:6328491
A;Accession: 146868
                                                                                                                                                                                                                              C;Species: Carausius morosus
C;Date: 30-Sep-1993 #text_change
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change
C;Accession: JC1416; S07157
R;Gaede, G.; Kellner, R.; Rinehart, K.L.; Proefke, M.L.
Biochem. Biophys. Res. Commun. 189, 1303-1309, 1992
Biochem. Biophys. Res. Commun. 189, 1303-1309, 1992
A;Title: A tryptophan-substituted member of the AKH/RPCH family
A;Reference number: JC1416; MUID:93129188; PMID:1482345
A;Accession: JC1416
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submitted to the Protein Sequence Data
submitted to The Proteins of kidney
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A; Residues: 1-10 <GAE1>
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A; Accession: G58502
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C;Species: unidentified bacterium
C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_change 10-Jul-1998
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Pred. No. 2.8e+05;
4; Mismatches 1
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1; Mismatches 1
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N,Alternate 1
C;Species: E:
C;Date: 30-JI
C;Accession:
R;Gaede, G.;
                                                                                   amine oxidase (copper-containing) (EC 1.4.3.6) - pig (fragment)
C;Species: Sus scrofa domestica (domestic pig)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 12-Oct-1989 #sequence_revision 31-Dec-1993 #text_change 06-Sep-1996
C;Accession: A32428
R;Van Ger Meer, R.A.; van Wassenaar, P.D.; van Brouwershaven, J.H.; Duine, J
Biochem. Biophys. Res. Commun. 159, 726-733, 1989
A;Title: primary structure of a pyrroloquinoline quinone (PQQ) containing pe
A;Reference number: A32428; MUID:89193662; PMID:2539124
A;Accession: A32428 a;MUID:89193662; PMID:2539124
A;Accession: A32428 a;MUID:89193662; PMID:2539124
A;Accession: A32428 a;MUID:89193662; PMID:2539124
A;Molecula type: protein
A;Note: the modified residue thought by the authors to be pyrroloquinoline q
C;Keywords: oxidoreductase; quinoprotein; topaquinone
C;Keywords: oxidoreductase; quinoprotein; topaquinone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A,Title: Primary structures of hypertrahalosaemic neuropeptides isolated from the entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom A; Reference number: $08995; MUID:90253659; PMID:2340112
A,Accession: $09138
A,Molecule type: protein
A; Residues: 1-10 GAE-
A,Note: the amino-terminal residue forms pyrrolidone carboxylic acid; therefore, A,Note: the amino-terminal residue formone; neuropeptide; pyroglutamic acid C; Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid F; L/Modified site: pyrrolidone carboxylic acid (Oln) #status experimental F; 10/Modified site: amidated carboxyl end (Thr) #status experimental
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A;Residues: 'Z',2-10 <GAB2>
C;Comment: Hypertrehalosemic factor II lacks the tryptophan modification.
C;Comment: This peptide raises hemolymph levels of trehalose in the cockr
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; glycoprotein; hormone; neuropeptide; p
F;I/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Binding site: carbohydrate (Trp) (covalent) #status experimental
F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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hypertrehalosemic hormone II -
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Chem. Hoppe-Seyler 371,
                          Local
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33.3%; Pred. No. 4.8e+03;
7. Mismatches 4;
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2; Mismatches 4
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ig heavy chain CDR3 region (clone 1-118B) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Accession: PT0231
R;Yamadd, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0231
A;Accession: PT0231
A;Molecule type: DNA
A;Residues: 1-9 cyAM-
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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A;Molecule type: protein
A;Residues: 1-7 <PHI>
Search completed: May 18, 2004, 10:05:40 Job time: 10.25 secs
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P810144
P82072
P82072
Q10582

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61
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ALL1 CYDPO
TT51 LITRU
ALL1 CYDPO
HTF1 PERAM
FAR8 MACRS
UF02 MOUSE
ALL1 CARMA
TKNC RANCA
TKNC PIG
ACST
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EFG_CLOPA
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   p49820 canis famil
p3103 proteus vul
p58707 anthopleura
p82158 cydia pomon
p82056 litoria rub
p81821 carcinus ma
p82152 cydia pomon
p04548 periplaneta
p83281 macrobrachi
p38640 mus musculu
p81822 carcinus ma
p22690 rana catesb
p01192 sus scrofa
p83146 bacteroides
p83146 bacteroides
p83176 carcinus ma
p22691 sus scrofa
p83194 bacteroides
p83176 carcinus ma
p22395 locusta mig
p80903 methanobact
p36414 pseudomonas
p25419 tenebrio mo
p21143 leucophaea
p1988 leucophaea
p05486 conus geogr
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Q03367;
Q1-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=cv. Lamuyo; TISSUE=Fruit, and Leaf;
MEDLINE=93099270; pubMed=1463853;
Kuntz M., Camara B., Weil J.-H., Schantz R.;
Wintz M., Camara B., Weil J.-H., Schantz R.;
Winte psbL gene from bell pepper (Capsicum annuum): plastid RNA editing also occurs in non-photosynthetic chromoplasts:";
Plant Mol. Biol. 20:1185-1186(1922).
-i- FUNCTION: This b-type cytochrome is tightly associated with the reaction center of photosystem II and possibly is part of the water-oxidation complex.
-i- SUBUNIT: Heterodimer of an alpha subunit and a beta subunit.
-i- SUBUNIT: Belongs to the psbE / psbF family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Capsicum annuum (Bell pepper).
Chloroplast.
Enderyota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots, asterids,
lamiids, Solanales, Solanaceae, Capsicum.
                                                                                                                                                                                                                                                                                                                         the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                         NON TER
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InterPro; IPRO6216; Cyt_b559.
PROSITE; PS00537; CYTOCHROME B559;
Chloroplast; Photosystem II; Heme;
                                                                                                                                         SEQUENCE
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10-OCT-2003
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4 (Rel. 29, Last sequence update)
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CORZ_PERAM
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PVKI_PERAM
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UKA2_HUMAN
UR2_FOLSP
REZI_LITRU
RE31_LITRU
ANG2_BOTJA
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Pred. No.
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                                                                                                          DB 1;
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                                                                                       2e+02;
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RESULT 2
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AC P36960;
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DE Lantibiotic
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CARNOBACTERIA;
CARNOBACTERIA;
CARNOBACTERIA;
RN NEDLINE=9232:
RN SEQUENCE
RY MEDLINE=9232:
RA Nes I.F.;
RT "PURTICAL
OC ARTIFICATION
CC ANCIDENCE
RY ADP1. ENVICO
CL -1- FUNCTION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIINE-89145002; PubMed-3226948; Gaede G., Hilbich C., Beyreuther K., Rineha "Sequence analyses of two neuropeptides of two he lubber grasshopper, Romalea microptera. Peptides 9:681-688(1988).
                                                                                                                                                                                                  Neuropeptide; Amidation; Flight;
MOD_RES 1 1 PYRI
MOD_RES 10 10 AMII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1990 (Rel. 16, Created)
01-FEB-1994 (Rel. 28, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
RO I (Hypertrehalossemin from the contraction)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P18110;
01-NOV-1990
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Carnobacterium
                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                      PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=7007;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Neoptera; Orthopteroidea; Orthop
Acridoidea; Romaleidae; Romalea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Romalea microptera (Lubber grasshopper)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Purification and characterization a Carnobacterium sp.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stoffels G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=92321768; PubMed=1622206;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=35782;
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                                                                                                                                                                                                                                                                                                                         InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SSUE=Corpora
                                                                                                                                                                                                                                                                                                                                             FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose major carbohydrate in the hemolymph of insects). SUBCIELLULAR LOCATION: Secreted. SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FUNCTION: Lanthionine-containing Active on Gram-positive bacteria. ibiotic; Bacteriocin; Lantibiotic. TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ROMMI
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEIQ 5
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                                                                                                                                                                                                                                                                                                                         IPR002047; AKH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Metazoa; Arthropoda; Hexapoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ٧
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(Rel. 29, Last sequence u)
(Rel. 40, Last annotation
.carnocin UI49 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA;
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                                                                                                                                                                    AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
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1163 MW;
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                                                                       29.5%;
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           opoda; Hexapoda; Insecta; Pterygota;
Orthoptera; Caelifera; Acridomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        741776D05B05B810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18;
Pred. No.
                                             Score 18; DB 1;
Pred. No. 1.1e+03;
2; Mismatches ;
                                                                                                                                                              jht; Pyrrolidone carboxylic a
PYRROLIDONE CARBOXYLIC ACID
AMIDATION.
056236745771A9C4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gudmundsdottir A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lart K.L. Jr.;
the AKH/RPCH-family
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CRC64;
                                                                                                    Length 10;
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                                          Gaps
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                                                   RESULT 5
HTF2_CARMO
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Best Local S
Matches 2
SEQUENCE.

SEQUENCE.

SPECIES=E, tiaratum; TISSUE=Corpora cardiaca;

SPECIES=E, tiaratum; TISSUE=Corpora cardiaca;

MEDLINE=90253559; PubMed=2340112;

Gaede G., Rinehart K.L. Jr.;

Gaede G., Rinehart K.L. Jr.;

"Primary structures of hypertrehalosaemic neuropeptides isolated from the corpora cardiaca of the cockroaches Leucophaea maderae, the corpora cardiaca of the cockroaches Leucophaea maderae, and the corpora cardiaca of the cockroaches Leucophaea maderae, and blatta orientalis
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30-MAY-2000
30-MAY-2000
                                                                                                                                                                                                                                                                                  Gaede G., Rinehart K.L. Jr.;
"Primary structure of the hypertrehalosaemic factor II
"Primary structure of the Indian stick insect, Carausius
determined by fast atom bombardment mass spectrometry."
Biol. Chem. Hoppe-Seyler 368:67-75(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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P11385;

P11385;

1 01-JUL-1989 (Rel. 11, Created)

T 01-FEB-1994 (Rel. 28, Last sequence update)

T 01-FCT-2003 (Rel. 42, Last annotation update)

AT 10-CCT-2003 (Rel. 42, Last annotation update)

E Hypertrehalosaemic factor II (HTF-II) (HRTH-II)

E Hypertrehalosaemic factor II (HTF-II) (HRTH-II)

Carausius morosus (Indian stick insect), and

Carausius morosus (Indian stick insect).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUB=Larva;

MEDLINE=98054539; PubMed=9392829;

Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., W
Davey M., East P.D., Thorpe A.;

"Lepidopteran peptides of the allatostatin superfam
Peptides 18:1301-1309(1997).

-!- SIMILARITY: Belongs to the allatostatin family.
SPECIES=C.morosus; TISSUE=Corpora MEDLINE=87157103; PubMed=3828078; Gaede G., Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Endopterygota; Lepidoptera; Glossata; Dit
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                              Heteronemiidae; Carausius.
NCBI_TaxID=7022, 7024;
                                                                                                                                          Extatosoma tiaratum (Stick insect).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Phasmatodea; Euphasmida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neuropeptide; Amidation.
MOD_RES 8 8
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                                                                                                                                                                                                                                                                                                                                                       STANDARD;
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40.0%;
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Pred. No. 1.4e+05
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                                         cardiaca;
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(HRTH-II)
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Phasmatoidea;
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MEDLINE-93129188; PubMed-1482345;
Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;
Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;
H. tryptophan.substituted member of the AKH/RPCH family isolated
a stick insect corpus cardiacum.";
Biochem. Biophys. Res. Commun. 189:1303-1309(1992).
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that
elevate the level of trehalose in the hemolymph (trehalose:
major carbohydrate in the hemolymph of insects).
-!- SUBCELULUAR LOCATION: Secreted.
-!- MASS SPECTROMETRY: MW-1308.61; METHOD-FAB.
-!- SIMILARITY: Belongs to the AKH / HXTH / RPCH family.
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28-FEB-2003
10-OCT-2003
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EI03_I
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Wabnitz P.A., Bowie J.H., Tyler M.J.,
"Peptides from the skin glands of the
Litori electrica. Comparison with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAREOHYDRATE-LINKAGE SITE.
SPECIES=C.morosus; TISSUE=Corpora cardiaca;
                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                      Aust
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata;
Amphibia; Batrachia; Anura; Ne
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Electrin
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PIR; S09138; S09138.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Litoria rubella (Desert tree frog)
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m bombardment
                                                                                                                                                                                                 SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Skin.
nibian defense peptide; Amidatio
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3; Conserv
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2; Conser
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(Rel.
(Rel.
                                                                                Conservative
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42,
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33.3%;
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                                                                                               24.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence
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                                                                           Score 15; DB 1;
Pred. No. 1.4e+05;
1; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Craniata; Vertebrata;
Neobatrachia; Hyloidea;
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Pred. No. 2.7e+03;
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                                                                                                                                                                                                                                                                                                          Wallace J.C.;
Australian buzzing tree frog
skin peptides from Litoria
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family isolated
                                                                                Indels
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Hylidae;
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RESULT

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CRC64

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RESULT
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Best Local S
Matches 2
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P35946;
01-JUN-1994
01-JUN-1994
10-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUSSO
                                                                                                                                                                                                                                               RNA polymerase beta sub-
fr. protein) (Fragment).
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NON_TER
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Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordari
Hypocremycetidae; Hypocreales; Nectriaceae; Nectria
NCBI_TaxID=70791;
                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstatton the European Bioinformatics Institute. There are no restrictions on its use by mon-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Verma J., Gangal S.V
Submitted (JUL-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P81010;
15-JUL-1998
Transferase; RNA-directed RNA
NON TER 10 10
SEQUENCE 10 AA; 1105 MW; 5
                                                                                                                                                                                                                                                                                                                                                   Phocine distemper virus (PDV).
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                    EMBL; D10371; BAA01208.1;
                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=11240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- ALLERGEN: Causes an allergic reaction
                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                             Gen. Virol. 73:1189-1194(1992).

FUNCTION: Probable component of the active polymerase. It may function in mRNA synthesis, capping, methylation and poly(A) synthesis of newly synthesized viral mRNAs, RNA editing of the gene transcript, and protein kinase activity.

CATALYTIC ACTIVITY: N nucleoside triphosphate = N diphosphate
                                                                                                                                                      SIMILARITY: Belongs to the paramyxoviruses L protein family.
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(Rel. 29,
(Rel. 42,
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5.V.;
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2, Last annotation updat
subunit (EC 2.7.7.48) (
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Pred. No. 1.4e
1; Mismatches
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P81350;
15-JUL-1998
15-JUL-1998
28-FEB-2003
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InterPro; IPRO00795; EF GTPbind.

PROSITE; PS00301; EFACTOR GTP; PARTIAL.

Elongation factor; Protein biosynthesis;

NON_TER 11 11
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P498Z0;
01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Flengsrud R., Skjeldal L.;
"Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5.";
Electrophoresis 19:802-806(1998).
-i- FUNCTION: This protein promotes the GTP-dependent translocation the nascent protein chain from the A-site to the P-site of the
                                                                                                                                                                                                                                                                   01-OCT-1996 (Rel. 34, Created)
15-JUL-1998 (Rel. 36, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
NADH-ubiquinone oxidoreductase 24 kDa
(EC 1.6.99.3) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                    CANFA
TISSUE=Heart;
MEDLINE=98163340; PubMed=9504812;
Dunn M.J., Corbett J.M., Wheeler C.I
"HSC-2DPAGE and the two-dimensional
dog heart proteins.";
Electrophoresis 18:2795-2802(1997).
                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              - -
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                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                Canis familiaris
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=1501;
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SIMILARITY: Belongs to
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98 (Rel. 36, Last :
03 (Rel. 41, Last :
n factor G (EF-G)
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.oa; Chordata; Craniata; Ver
.oa Carmivora; Fissipedia;
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annotation update
(CP 5) (Fragment)
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                                                                 C.H.;
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                                              electrophoresis database
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Canis.
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P-site of the
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Matches
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Best Local
                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no was modified and this statement is not removed. Usage by and for commercia entities requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@sisb-sib.ch).
                                                                                                                                                                                                            WEDLINE=3087186; PubMed=1454536; Smith M.D., Longo M., Gerard G.F., Chat """ and characterization of genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; L04163; AAA25660.1;
PIR; S35490; S35490.
REBASE; 1541; PVuI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUL-1993 (Rel 26, Creat
01-JUL-1993 (Rel 26, Last
10-OCT-2003 (Rel 42, Last
Type II restriction enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oxidoreductase; NAD; Ubiquinone; Mitochondrion; Iron-sulfur; Iron; 2Fe-2S.

NON_TER 11 11
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HSC-2DPAGE; P49820; DOG.
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            Restriction NON_TER
                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. STRAIN=ATCC 13315;
                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=585
                                                                                                                                                                                                                                                                                                                                                                                                                           Enterobacteriaceae;
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PROSITE; PS01099; COMPLEX1_24K; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Proteus vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: TRANSFER OF ELECTRONS FROM NADH TO THE RESPIRATORY CHAIN. THE IMMEDIATE ELECTRON ACCEPTOR FOR THE ENZYME IS BELL TO BE UBIQUINONE. COMPONENT OF THE FLAVOPROTEIN-SULFUR (FP) FRACMENT OF THE ENZYME.

CATALYTIC ACTIVITY: NADH + ubiquinon = NAD(+) + ubiquinol.

CATALYTIC ACTIVITY: NADH + acceptor = NAD(+) + reduced accept COFACTOR: Binds I 2Fe-2S cluster (Potential).

SUBUNIT: Mammalian complex I is composed of 45 different subusubcellular LOCATION: Matrix and cytoplasmic side of the mitochondrial inner membrane.

SIMILARITY: Belongs to the complex I 24 kDa subunit family.
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                           system;
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                             Hydrolase; Nuclease;
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annotation update)
PvuI (EC 3.1.21.4)
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Pred. No. 4.7e+03;
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                                                                                                                                                                                                                                                                                                                  Chatterjee D.K.; enes for the Pvul restriction
                                                                                                                    moved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
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                             Endonuclease
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Query Match Best Local S Matches

24.6%; Similarity 60.0%; 3; Conservative

Pred. No. 4.76 1; Mismatches Score 15; Pred. No.

4.7e+03;

DB 1;

Length 11;

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SEQUENCE

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Matches 2
Query Match
Best Local Similarity
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P82158;
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FIRN ANTEL STAN
P58707;
28-FEB-2003 (Rel. 4
.28-FEB-2003 (Rel. 4
10-CCT-2003 (Rel. 4
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MOD_RES
SEQUENCE 4 A
                                                                                                                                       MOD RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidaa; Tortricidae; Olethreutinae; Cydia.
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Eukaryota; Metazoa; Chidaria; Anthozoa;
Nynantheae; Actiniidae; Anthopleura.
NCBI_TaxID=6110;
                                                                                                                                                                                             Duve H., Johnsen A.H., Maestro J.-L., Scott i Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin supertides 18:1301-1309(1997).
-!- SIMILARITY: Belongs to the allatostatin Neuropeptide; Amidation.

Neuropeptide; Amidation.

MOD RES 7 AMIDATION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE.
TISSUE=Larva;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=82600;
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Proc. Natl. Acad.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anemone neuropeptide containing an unusual amino-terminal blocking
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=98054539; PubMed=9392829;
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:- TISSUE SPECIFICITY: Neuron specific.
:- MASS SPECTROMETRY: NW=549.3; METHOD=FAB
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nilarity 50.0%;
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. 39, Last annotation.
                                                                                                                                            873 MW;
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                                                                                                                                                                                                                                                                                                                                                                 the allatostatin superfamily.";
                                                                                                                                       AMIDATION.
672879CABB569350
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Pred. No. 1.4e+05;
1; Mismatches 1
    Score
Pred.
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Best Local S
Matches 2
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P81821;
30-MAY-2000
30-MAY-2000
30-MAY-2000
    "Isolation and identification of multiple neuropeptides allatostatin superfamily in the shore crab Carcinus maes Eur. J. Biochem. 250:727-734(1997).

-I- FUNCTION: May act as a neurotransmitter or neuromodule.
-I- SIMILARITY: Belongs to the allatostatin family.

Neuropeptide; Amidation; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                                                                                           TISSUE=Cerebral ganglion, and Thoracic ganglion; MEDLINE=98121193; PubMed=9461295; Duve H., Johnsen A.H., Maestro J.-L., Scott A.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VCBI_TaxID=6759;
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STANDARD;

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39, Created)
39, Last sequences
39, Last annotations

Last sequence update)
Last annotation update)

(Common shore crab) (Green crab)

Brachyura;

Scott A.G.,

Jaros P.P.,

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Matches
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P82065;
28-FEB-2003 (Rel. 4
28-FEB-2003 (Rel. 4
10-OCT-2003 (Rel. 4
                                                                                                                                               MOD RES
MOD RES
SEQUENCE
                                                                                                                                                                                                                   Tyler M.J., Wallace J.C.;

"The structure of new peptides from the Australin red tree frog

"Litoria rubella". The skin peptide profile as a probe for the study

of evolutionary trends of amphibians.";

Aust. J. Chem. 49:955-963(1996).

-!- FUNCTION: May act as a neuromodulator or neurotransmitter.

-!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECTROMETRY: May-965; METHOD-FAB.

-!- MASS SPECTROMETRY: May-965; METHOD-FAB.

Amphibian defense peptide; Amidation; Neuropeptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tryptophyllin 5.1.
Litoria rubella (Desert tree frog).
Litoria rubella (Desert tree frog).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota, Metazoa; Chordata; Craniata; Hyloidea;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Skin secretion;
Steinborner S.T., Wabnitz P.A., Waugh R.J.,
                                                                                                                                                                                                     Pyrrolidone carboxylic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE,
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                                 4 EIQFMH 9
                                                                     Similarity
2; Conserve
QIPWFH 6
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                                                                       Conservative
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41, Last sequence update)
42, Last annotation update)
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Pred. No. 1.4e
2; Mismatches
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                                                                 DB 1; L.
1.4e+05;
2;
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Hylidae;
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Perfect score:
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1: sp_archea:*

2: sp_bacteria:*

3: sp_fungi:*

4: sp_human:*

5: sp_invertebrate

6: sp_manmal:*

7: sp_mhc:*

10: sp_phage:*

11: sp_rodent:*

12: sp_virus:*

13: sp_vertebrate

14: sp_unclassifi

15: sp_bacteriap:*

16: sp_bacteriap:*
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Match
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125.164 Million cell updates/sec
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88888779999777777
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sp_plant: *
sp_plant: *
sp_virus: *
sp_vertebrate: *
sp_vertebrate: *
sp_vertebrate: *
sp_bacteriap: *
sp_archeap: *
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Copyright (c) 1993 - 2004 Compugen Ltd.
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sp_human:*
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P82081
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                                                    O9ps71 agkistrodon
O9t178 canis famil
O92766 canine dist
071066 canine dist
P82081 limnodynast
O13591 saccharomyc
O15342 homo sapien
O8mesO anoda crist
O85631 avian carroi
O23876 dictyosteli
O23874 paramecium
O28742 oryctolagus
O855j20 gallus gall
  Q05403 saccharomyc
Q40659 oryza sativ
P82082 limnodynast
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  RESULT
Q9TT78
ID Q9
AC Q9
DT 01
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Matches 4
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Q9PS71;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UNN-2020 (TrEMBLrel. 21, Last annotation update)
01-UNN-2002 (TrEMBLrel. 21, Last annotation update)
Fibrinolytic metalloproteinase (Fragment).
Agkistrodon contortrix.
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Agkistrodon.
MCBI TaxID=8720;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=91378546; PubMed=1898066; Guan A.L., Retzios A.D., Henderson G.N., Markland F.S.Jr.; Guar A.L., Retzios A.D., Henderson G.N., Markland F.S.Jr.; "Purification and characterization of a fibrinolytic enzyme of the southern copperhead snake (Agkistrodon contortrix contortrix)."; Arch. Biochem. Biophys. 289:197-207(1991).

NON TER 11
11
SEQUENCE 11 AA; 1209 MW; 7CA02D1D41E8772B CRC64;
  Q9TT78
Q9TT78;
Q1-MAY-2000 (TrEMBLrel. 13,
Q1-MAY-2000 (TrEMBLrel. 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            EQUENCE
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                                                                                                                                                                                                                              Similarity 4; Conserv
                                                                                                                                                     HNLG 10
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Q8RKU3
Q83928
Q43928
Q44001
Q9FXL0
Q9GD68
Q9GD68
Created)
Last sequence update)
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Q44468 aeromonas v
Q874468 aeromonas p
Q87401 borreila bu
Q43928 aeromonas p
Q44001 aeromonas e
Q96x10 lilium long
Q96x10 lilium long
Q96x10 lilium long
Q96x10 limnodynast
Q15894 homo sapien
Q34909 locusta mig
Q90493 eopsaltria
Q43960 azotobacter
Q15891 homo sapien
Q8366 glycine max
Q916x10 hordeum vul
Q910x10 hordeum vul
Q910x10 hordeum vul
Q910x10 pachola angu
Q90x10 pichia angu
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Q90x10 pan paniscu
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from venom

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Gaps

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O92766
ID O92766;
AC O92766;
AC O92766;
DT O1-NOV-1998
DT O1-NOV-1998
DT O1-NOV-1998
OS Canine distered oc Viruses; ssrr
OC Paramyxovirio
OX NCBI TaxID=11
RN [1]
RP SEQUENCE FROM
RP STRAIN=Dog ##
RA Liermann H.,
RT "Genetic anal
RT proximal coud
RT morbillivricud
RI SUBmitted (SI
RM SEQUENCE 9
SQUENCE 9
RESULT
O71066
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071066;
01-AUG-1998
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Fusion prote
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Thymidylate
TS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Genetic analysis of the central untranslated genome region proximal coding part of the F gene of wild-type and vaccine morbilliviruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; AF026237; AAC09167.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canine distemper virus.
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
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                 Canine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=Dog #5526/89;
Liermann H., Harder T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL;
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  Viruses;
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distemper virus.
s; ssRNA negative-strand
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-1998 (TremBirel. 07,
-2001 (TremBirel. 19,
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ba; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                             1011 MW;
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                                                                                                                                                                                                                                                                                                                                                   31.1%; 5-
100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6731A1E059CAA867 CRC64;
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Last annotation update)
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Pred. No. 1e+0
1; Mismatches
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Pred. No.
                                                                                                                                                                      PRT;
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  viruses;
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1e+06;
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  Mononegavirales,
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Canis.
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O13591;

O1-JAN-1998 (TrEMBLrel. 05, 0

O1-NOV-1999 (TrEMBLrel. 12, 1

O1-OCT-2002 (TrEMBLrel. 22, 1

ORF YNL337W (Fragment).
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P82081;
01-MAY-2000
01-MAY-2000
01-MAY-2000
DYNASTIN 3.
Obermaier
Submitted
[2]
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                                                            SEQUENCE FROM N.A.
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TISSUE=TIBIAL GLAND;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=104894;
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NCBI_TaxID=11232
                                                                                                  NCBI_TaxID=4932;
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                      B., Piravandi
(MAY-1996) to
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100.0%; Pr
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the
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Last
                      Rinke M.;
EMBL/GenBank/DDBJ
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SEQUENCE FROM N.A.
STRAIN=Dog #10757/96;
Liermann H., Harder T., Haas L.;
Liermann H., Harder T., Haas L.;
"Genetic analysis of the central untranslated genome region proximal coding part of the F gene of wild-type and vaccine morbilliviruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetales; Saccharomycetaceae; Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Raftery M.J., Bradford A.M., Bowie J.H., Wallace J.C., Tyler M.J.; "Peptides from Australian frogs. The structure of the dynastins from the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Limnodynastes terraereginae (Northen banjo frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachi
Limnodynastinae; Limnodynastes.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Chem. 46:833-842(1993).
SPECTROMETRY: MW=1236; METHOD=FAB.
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Aml1 protein
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Q8MES0;
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01-JUN-2003;
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EMBL; Z716;
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SEQUENCE
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MEDLINE=96226397; PubMed=8634147;

Levanon D., Bernstein Y., Negreanu V., Ghozi M.C., Bar-Am I.,

Levanon D., Bernstein Y., Officer Y.;

Aloya R., Goldenberg D., Lotem J., Groner Y.;

"A large variety of alternatively spliced and differentially express

menNas are encoded by the human acute myeloid leukemia gene AML1.";

DNA Cell Biol. 15:175-185(1996).

EMBL; X90978; CAA62465.1; -.
Pfeil B.E., Brubaker C.L., Craven "Phylogeny of Hibiscus and the tri chloroplast DNA sequences of ndhF Syst. Bot. 27:333-350(2002).
EMBL; AF384567; AAM50405.1; -. GO; GO:0009507; C:chloroplast; IEA
                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eneurosids II; Malvales; Malvaceae; Malvoideae; Anoda.
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Mammalia; Eutheria;
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                                        Craven L.A., Crisp M.D.;
the tribe Hibisceae (Malvaceae)
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"3'-terminal region of avian carcinoma virus MH2 she
"lements with avian sarcoma viruses Y73 and SR-A.";
J. Virol. 52:703-705(1984).
BMBL, K03100; AAA42388.1;
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Viruses; Retroid viruses; Retroviridae; Alpharetrovirus.
NCBI_TaxID=11958;
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Eukaryota; Mycetozoa; Dictyosteliida;
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Q35374;
01-NOV-1996
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Q28742;
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01-JUN-2003
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STRAIN=stock 172;
STRAIN=stock 172;
MEDLINE=87055241; PubMed=3023187;
MEDLINE=87055241; PubMed=3023187;
MEDLINE=67055241; PubMed=3023187;
MEDLINE=6705241; PubMed=5705241; PubMed=5705241; PubMed=5705241; PubMed=57052418; PubMed=5705241; PubMed=5705241; PubMed=5705241; PubMed=5705241; PubMed=5705241; PubMed=5705241; PubMed=5705241; PubMed=57052
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NON TER
SEQUENCE
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replication origin.",
Gene 44:243-253(1986).
EMBL; M15280; AAA79267.1; -.
GG; GG:0005739; C:mitochondrion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Lagomorpha; Leporidae; NCBL TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alpha-myosin heavy chain (Fragment).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                    ventricular myosin heavy chains.";
Proc. Natl. Acad. Sci. U.S.A. 81:3044-3048(1984)
EMBL; K01698; AAA31415.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE=84221901; PubMed=6328491;
Friedman D.J., Umeda P.K., Sinha A.N
Rabinowitz M.;
"Characterization of genomic clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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3; Conserv
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                                                         QKMHD 5
                                                                                                                  QFMHN 10
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11 AA;
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(TrEMBLrel. 24, Last annotation update)
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Pred. No. 1.5e
3; Mismatches
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Pred. No. 1e+06;
1; Mismatches
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.1.5e+04;
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Best Local S
Matches
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Best Local S
Matches 2
Q40659;
Q40659;
01-NOV-1996
01-NOV-1996
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Gallus gallus (Chicken).
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Meognathae; Galliformes; Phasianidae; Phasiani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q8JJ20 PRELIMINARY; PRT; 7 AA. Q8JJ20; Q9JJ20; Created) 01-OCT-2002 (TrEMBLrel. 22, Last sequence update) 01-OCT-2002 (TrEMBLrel. 22, Last annotation update) Extracellular fatty acid binding protein (Fragment)
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01-NOV-1996 (TrEMBLrel. 21, Last sequence update)
01-NOV-1996 (TrEMBLrel. 22, Last annotation update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
DNA for ORF'S from chromosome XV (Fragment).
Saccharomyces cerevisiae (Baker's yeast).
Eukaryota, Fungi, Ascomycota, Saccharomycotina, Saccharomycetes,
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zumstein E., Pearson B.M., Kalogeropoulos A., Schweizer M.; "A 29.425 kb segment on the left arm of yeast chromosome XV more than twice as many unknown as known open reading frame Yeast 11:975-986(1995).
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NON TER 1 1
SEQUENCE 7 AA; 780 MW; 72CI
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"Cloning and sequencing of 3'
Submitted (FEB-2002) to the EL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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STRAIN=FY1679;
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the EMBL/GenBank/DDBJ databases
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DE Alpha-amylase (Fragment).

OS Oryza sativa (Rice).

OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

OC Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

OC Ehrhartoideae; Oryzeae; Oryza.

OX NCBI TaxID=4530;

RN [1] - RXID=4530;

RN SEQUENCE FROM N.A.

RX MEDLINE=91078641; PubMed=2258052;

RX MEDLINE=91078641; PubMed=2258052;

RX Kumagai M.H., Shah M., Terashima M., Vrkljan Z., Whitaker J.R.,

RA Rodriquez R.L.;

RT cerevisiae.";

RT cerevisiae.";

Gene 94:209-216(1990).

DR Gramene; Q40659; -.

BR Gramene; Q40659; -.

BR NON TER 8 AA; 948 MW; EBC694444732D6D6 CRC64;

Query Match

Best Local Similarity 28.6%; Pred. No. 1e+06;

Matches 2; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Matches 2; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Search completed: May 18, 2004, 10:04:47

Job time: 30.25 secs
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Gapop 10.0 , Gapext 0.5
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Parathyroid hormone antigenic peptide 2-12 11-FEB-2003 ABG72607; ABG72607 standard; peptide; 11 (first entry) A

ALIGNMENTS

Human; parathyroid hormone; antigen; hypercalcaemia; osteoporosis; primary hyperparathyroidism; mouse; rat; bovine; porcine; canine.

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RESULT 1
ABG72607
ID ABG72607
XX ABG7
XX ABG7
XX ABG7
XX Para
XX Huma
XX Huma
XX Homc
OS Mus
OS Sus
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Rattus sp.
Bos taurus.
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(ZAHR/) ZAHRADNIK R J. (LAVI/) LAVIGNE J R.

Zahradnik RJ, Lavigne 占

WPI; 2003-066685/06.

New parathyroid hormone (PTH) antigenic peptide inducing the and isolation of antibodies having an affinity to it, useful determining bioactive PTH levels in serum, plasma and/or cell media. formation for loulture

Claim 1; Page 5; 11pp; English

AAR91646 AAB07467 AAB86221

The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, being

WPI; 2003-066685/06

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC formed from the N-terminus of parathyroid hormone (PTH). Also included CC are, (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one first peptide antigen comprising a peptide fragment of PTH; (b) candinistering the first peptide antigen to a host animal to induce antibody production; (c) monitoring the antibody titre produced; (d) contisted antisers produced in the host animal; and (e) selecting antisers produced in the host animal; and (e) selecting contisted antisers produced in the host that is capable of binding to a second peptide antigen; (2) an antibody (ab) produced by the method; and (3) test kits and analytical procedures used for the compositions of the present invention are useful for determining compositions of the present invention are useful for determining compositions of the present invention are useful for determining compositions of the present invention are useful for determining compositions and methods of the present invention, as creater affinity for PTH, and in particular, are designed to have a novel crecognition for amino acid residues extending beyond the first N-terminal compositions of prince prince prince are an important parameter in patients suffering from hypercalcaemia, osteoporosis and primary compositions and primary and composition conservating amino acids 2-12 of human, mouse, rat, porcine, consider and bovine PTH
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Matches
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                                                                                                                                                                              US2002110871-A:
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                            (ZAHR/) ZAHRADNIK R J.
(LAVI/) LAVIGNE J R.
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Sequence

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The invention relates to a new antigenic peptide for inducing the commation and isolation of antibodies having an affinity to it, being commed from the N-terminus of parathyroid hormone (PTH). Also included care, (1) a method for producing antibodies useful in the determination of comprising the first peptide antigen comprising: (a) providing at least one cadministering the first peptide antigen to a host animal to induce antibody production; (c) monitoring the antibody titre produced; (d) compares from the isolated antigen to a host animal to induce antipers from the isolated antigen produced in the host animal, and (e) selecting cantisers produced in the host animal; and (e) selecting compares from the isolated antigen; (2) an antibody (ab) produced by the method; and (3) test kits and analytical procedures used for the comparistions of the present invention are useful for determining the antigens; antibodies and methods of the present invention, as compared to prior art, have the particular advantages of possessing greater affinity for PTH, and in particular, are designed to have a novel crecognition for amino acid residues extending beyond the first N-terminal CC presidue, and further have negligible cross-reactivity with the large compared to prior art, have the particular, are designed to have a novel crecognition for animo acid residues are an important parameter in CC paptide representing amino acids 1-12 of human, mouse, rat, porcine, cc canine and bovine PTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New parathyroid hormone (PTH) antigenic peptide inducing the formation and isolation of antibodies having an affinity to it, useful for determining bioactive PTH levels in serum, plasma and/or cell culture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2;
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RESULT 3
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Best Local S
Matches 10
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                                                                                                                                                                                               AAY9696B;
(GARD/)
(KRON/)
(POTT/)
                                                                                                             Homo sapiens
                                                                                                                                                          Parathyroid hormone N-terminal signaling domain (residues 1-11).
                                                                                                                                                                                                                 AAY96968 standard; peptide;
                                     31-DEC-1998;
                                                       30-DEC-1999;
                                                                                           WO200039278-A2
                                                                                                                              parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tether1; osteoporosis.
) GARDELLA T J.
) KRONENBERG H M.
) POTTS J T.
                                                                                                                                                                                                                                                                                                    l Similarity
10; Conser
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                                      98US-0114577P
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90.9%;
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Zahradnik RJ,

Lavigne JR

Gardella

, LT

Kronenberg

HM,

Potts JT,

Jueppner H;

JUEPPNER

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Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of pTH(1-34) or a pTH-related protein (PTH-TP) (1-34); R is the PTH-1 receptor signal sequence; and R is (a portion of) pTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (Claimed). The new compound can be administered by inhalation unlike the large native PTH or PTH-P which avoids the need for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               regular injections to treat osteoporosis
                                                  New antigenic peptide(s) fr
generated using them, able
                                                                                                                                                                                                                                       04-APR-1996
                                                                                                                                                                                                                                                                                                                       diagnosis;
                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR91644 standard;
                                                                                                                                                                                                          28-SEP-1994;
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                                                                                                                                                                                                                                                                                            Synthetic
                                                                                             1996-180391/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Page 92; 119pp;
                                                                                                                                                                                                                                                                                                                         parathyroid hormone; 
psis; active hPTH 1-37
                                                                                                                                                                                                                                                                                                                                                               parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
9; Conserv
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                                                                                                                          Forssmann
                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                               94DE-04434551
                                                                                                                                                                                                            94DE-04434551
                                                                                                                                                                                                                                                                                                                                      hormone; hPTH;
                                                                                                                                                                                                                                                                                                                                                               hormone antigenic peptide hPTH 1-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75.4%;
81.8%;
                                                     from
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Pred. No.
                                                 human parathyrold normone - and distinguish between active and
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                                                                                                                                                                                                                                                                                                                                      antigen; alpha-helix; immunogen;
                                                                   parathyroid hormone - and antibodies
                                                                                                                          Maegerlein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.059;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 3;
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2

Page 4;

German

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Matches

Similarity

73.8%;

Score 45; Pred. No.

Length 10

Mismatches 0 DB 3; .082;

Indels

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Conservative

SVSEIQFMHN 10 SVSEIQLMHN

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Sequence

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Best Local :
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                                   The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipophilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin firetree.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Parathyroid hormone; lipolysis; cosmetic; dermatological; skin care, slimming treatment; cellulite; skin firming.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY68767;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY68767 standard;
                                                                                                                                                                                               in topically applied cosmetic composexcessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                   07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                               27-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amino acids 1-10 of a parathyroid hormone (pTH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-MAY-2000
                                                                                                                                                                  Claim 1; Page 8; 18pp; French.
                                                                                                                                                                                                                           New parathyroid hormone fragment peptides,
                                                                                                                                                                                                                                                       WPI; 2000-171243/15.
                                                                                                                                                                                                                                                                                  Lintner
                                                                                                                                                                                                                                                                                                            (SEDE-) SEDERMA
                                                                                                                                                                                                                                                                                                                                        17-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9;
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                                                                                                                                                                                                                                                                                                                                                                  99WO-FR001687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 73.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10
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Pred. No. 0.082;
D; Mismatches
                                                                                                                                                                                                               compositions
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                                                                                                                                                                                                             used as lipolysis stimulants for slimming treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 10;
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AAB86219
RESULT 7
ABR44166
ID ABR4
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AC ABR4
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                                                                                                                                                                                                                                                                                   This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CC of diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or CC typer-parathyroidism. The method (unlike known assays) recognizes that CC (i) some fragments of (A) shorter than the complete (84 aa) peptide are CC cinactive, and (ii) that apparently intact peptide may be biologically CC are antagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of CC conditions of the provides a true measure of the content of complete represents a peptide fragment used to illustrate the method of the other represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                    Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 3; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-376318/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Armbruster
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                                            ABR44166 standard; peptide; 10
                                                                                                                                                                                                                                             Sequence 10
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                                                                                                                                                                                    Similarity
9; Conserv
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                                                                                                                        SVSEIQLMHN
                                                                                                                                                                                                                                                 AA;
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                                                                                                                        10
                                                                                                                                                                                                 73.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunogenic peptide
                                                                                                                                                                                    Score 45; DB
Pred. No. 0.08
0; Mismatches
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ABR44166

17-DEC-1999;

99DE-01061350

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RESULT 8
AAB86225
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a fusion peptide (Tat-hPTHDP), where self cell penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmetic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endotheliu does not cause skin disease and has superior lipolysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fusion paptide; tat; hPTHDP; parathyroid hormone; skin; cosmetic; lipolysis; human; hPTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-SEP-2001; 2001KR-00060245
15-MAR-2002; 2002KR-00014062
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human
                                                                                                       Parathyroid hormone; hPTH; immunogenic peptide; human; epitopediagnosis; calcium-metabolism disorder; osteopathy; antagonist, hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                       Human parathyroid hormone immunogenic peptide SEQ ID
                                                                                                                                                                                                          AAB86225;
                                                                                                                                                                                                                                AAB86225 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 6; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel fusion peptide comprising self cell-penetrating Tat peptide bound to human parathyroid hormone-derived peptide, useful as component of skin slimming cosmetic composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GLDS ) LG HOUSEHOLD & HEALTH CARE LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2002; 2002WO-KR000835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003035697-A1
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                                                          DE19961350-A1
                                                                               Homo sapiens
                                                                                                                                                                                  03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10 AA;
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                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                             73.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (hPTH) fragment (residues 1-10)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cho w,
                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                             Score 45; DB 6;
Pred. No. 0.082;
                                                                                                                                                                                                                                                                                                                                                 Mismatches
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RESULT 9
AAR91645
ID AAR9
XX AAR9
AC AAR9
XX Huma
XX Huma
XX Huma
XX Huma
XX Synt
XX Synt
XX OA-1
PN DE44
PN DE44
PN 28-4
PR 28-4
XX AG
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 9
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                                                                                                                                                                                                                                                                                                                                04-APR-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human parathyroid hormone; hPTH; antigen; alpha-helix; diagnosis; active hPTH 1-37.
                                  WPI; 1996-180391/19
                                                                                                                                                                                                                                                                        28-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                            DE4434551-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human parathyroid hormone antigenic peptide hPTH 1-9.
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8; Conserv
                                                                                                                                                       FORSSMANN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VSEIQFMHN 10
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                                                                                           Forssmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                           Σ,
                                                                                              Hock
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Pred. No. 1.4e
0; Mismatches
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                                                                                           Maegerlein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .4e+06;
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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a specific example of claimed immunogenic peptides having a sequence from hPTH(1-37) which includes the N- or C-terminal alpha-helical region and/or the non-structured region of the hormone. Antibodies and their binding fragments generated by injecting animal with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antigenic peptide(s) from human parathyroid hormone - and antibodies generated using them, able to distinguish between active and inactive forms of the hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GARD/)
(KRON/)
(POTT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                    2000-452384/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            parathyroid reformation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ) KRONENBERG H M.
) POTTS J T.
JIPPDOVITE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JUEPPNER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kronenberg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hormone; N-t; resorption;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-terminal; signaling ion; remodeling; tether
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Pred. No. 1.4e
0; Mismatches
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1.4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jueppner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            naling domain;
tether1; oste
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Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH), L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of pTH(1-34) or a pTH-related protein (PTH-rp) (1-34); R_1 is tip PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing CAU in a mammalian cell having PTH-1 receptors, or screening for a peptide of the sequence of th

binding R_1 is the

(where

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

Claim

Page 93; 119pp;

English.

0,

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Query Match
                    Sequence 9
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63 .99

Score

39,

DB 4.

Length

This invention describes a novel method for the preparation of peptides CC (II) that contain both cyclic and linear peptide fragments comprises CC sequential reaction of a resin-bound linear fragment with the cyclic fragment in N-protected form and optionally other linear fragments. The CC bind to hPTH receptors and act as agonists or antagonists of hPTH. The CC method is particularly used to prepare cyclic peptide analogs of CC parathyroid hormone (PTE) or PTH-related peptides which are useful for CC treating diseases that respond to treatment with agents that bind to PTH CC receptors (with or without activation of adenylyl cyclase activity), e.g. CC hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, osteopenia, hyper- or hypo-c

Disclosure; Page 75;

85pp; English.

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RESULT 11
AAB86
XX AAB86
XX AAB86
XX Parat
XX Pa
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                                                                                    This invention describes a novel method for determining (M1) the content CC of active parathyroid hormone (A) by treating a sample with (i) antibody CR (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CR recognizes an epitope within the receptor-binding site of (A). The number of modecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or typer-parathyroidism. The method (unlike known assays) recognizes that (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically cinactive, and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-cerminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hyper-parathyroidism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTHrP which avoids the nu regular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibod reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
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                                                                   invention
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Pred. No. 1.4e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Convergent synthesis of peptides for treating e.g. bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sledeski
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The invention relates to a novel parathyroid hormone (PTH) peptide (ABB01859) and parathyroid hormone-related peptide (PTHrP, ABB01860), and biologically active derivatives thereof (ABB01857-B01858, ABB01861). The peptides of the invention are at least 85% identical to the generic peptide of the formula: X1-Val-Ser-Glu-X2-Gln-Leu-X3-H15-X4-X5-CGly-Lys-X6 (ABB01857), where: X1 is Ser or Ala; X2 is 11e or Lys; X3 is Ser or Ala; X2 is 11e or Lys; X3 is Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; provided that the peptide is not PTHrP(1-4). The peptides of the invention also encompass fragments of peptides of the invention consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-terminal derivatives. PTH is a major regulator of calcium homeostasis, and is necessary for the normal function of the gastrointestinal, seletal, neurological system, neuromuscular and cardiovascular systems. It binds to both PTH-1 receptors on osceoblasts and renal tubular cells, and to the recently identified PTH-2 receptor. PTH has a potent anabolic effect on the skeleton, and mediates calcium rebsorption, enhances phosphate clearance and vitamin D synthesis in the kidney. A homologous calls of protein hormones, the PTH-related proteins (PTHrP) minute some of the renal and skeletal actions of PTH, and also bind to the PTH-1
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Matches
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11
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                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 26;
                                                                                                                                                                                                                                                                                                                                                                                            Parathyroid hormone
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                                                                                                                                                                                                                                                                                                                                                                            hyroid hormone (PTH) peptides, PTH related peptides that encode them, useful for treating osteoporosis.
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KRONENBERG H M.
POTTS J T.
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JUEPPNER H.
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CX PTH-1/PTH-2 receptor agonists
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POTTS J T.
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7; Conserv
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1.4e+06;
ches 1;
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compounds of the structure or formula S-(L)_n-B, R 1-S-(L)_n-R or S-(L)_R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R 1 is terminal sequence; and R is (a portion of) PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases

binding R 1 is the

(where

s-(L)_n

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

WPI; 2000-452384/39

Claim 4; Page 92; 119pp; English.

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RESULT 15
AAB01863
The invention relates to a novel parathyroid hormone (PTH) peptide (CAB801859) and parathyroid hormone-related peptide (PTH-P, AAB01860), and Cobiologically active derivatives thereof (AAB01857-B01858, AAB01861-CC B01869). The peptides of the invention are at least 85% identical to the generic peptide of the formula: X1-Val-Ser-Gin-Y2-Gin-Lea-X3-His-X4-X5-CC generic peptide of the formula: X1-Val-Ser-Gin-Y2-Gin-Lea-X3-His-X4-X5-CC gry-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is CC with the peptide is not PTH-P(1-14). The peptides of the CC invention also encompass fragments of peptides of the invention consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-CC errainal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal, cCC and is necessary for the normal function of the gastrointestinal, cCC it binds to both PTH-1 receptors on osteoblaste and renal tubular cells, and to the recently identified PTH-2 receptor. PTH has a potent anabolic effect on the skeleton, and mediates calcium reabsorption, enhances compassed to the clearance and vitamin D synthesis in the kidney. A homologous
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTHrP which avoids the need for regular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthesis; bone synthesis; agonist; osteoporosis; non-parenteral delivery.
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                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 26; 73pp; English
                                                                                                                                                                                                                                                                                                                                                                          Parathyroid hormone (PTH) peptides, acids that encode them, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-339693/29
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) POITS J T.
JUEPPMTT
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Pred. No. 1.4e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jueppner H;
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calls of protein hormones, the PTH-related proteins (PTHrP) mimic some of the renal and skeletal actions of PTH, and also bind to the PTH-1 creeptor. They do not bind to the PTH-2 receptor. The peptides of the civention are either agonists of PTH-1 and PTH-2 receptors (AAB01858, CC AAB01861-B01866) or are PTH-1/PTH-2 receptor antagonists (AAB01867-CC B01869). PTH-1/PTH-2 receptor agonists are useful for the treatment of conditions characterised by a decrease in bone mass, such as costeoporosis. PTH-1/PTH-2 receptor antagonists are useful for treating cc medical disorders that arise from excessive or altered action of the PTH-CC medical condition. Detectably labelled peptides of the invention are also useful in the determination of rates of bone formation, bone resorption consentional synchetic chemistry, and can be delivered to a patient via conventional synchetic chemistry, and can be delivered to a patient via conventional receptor agonists
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Sequence 9 AA;

Matches Query Match Best Local 6 Similarity Conservative 54.18; 23 Score 33; DB 3; Pred. No. 1.4e+06; 2; Mismatches 1 Length 9; Indels 0, Gaps 0

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AVSEIQLLH 9

Search completed: May 18, 2004, 10:01:54 Job time : 46.75 secs

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Minimum DB
Maximum DB
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Listing first 45 summaries
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd
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9 US-09-730-174A-6
9 US-09-730-174A-5
9 US-09-730-174A-2
9 US-09-730-174A-3
9 US-09-730-174A-3
1 US-09-730-174A-1
1 US-10-168-185-7
4 US-10-168-185-7
4 US-10-168-185-7
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1 US-10-192-673-7
1 US-10-192-673-7
1 US-10-033-741-61
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Sequence 4, Appli
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Sequence 61, Appli
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Patent No. US20020110871A1
Patent No. US20020110871A1
Patent No. US20020110871A1

GENERAL INFORMATION:
APPLICATE: Zahradnik, R.J.
FITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho FILE REFERENCE: INUNECOLA
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 4
LENGTH: 12
TYPE: PRI
ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Induces Formation of Antibodies and Isolates said Antibodies
US-09-730-174A-4
RESULT 2
US-09-730-174A-6
US-09-730-174A-6
US-09-730-174A-6
J. Patent No. US20020110871A1
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid F, FILE REFERENCE: IMUNE-001A
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US-09-730-174A-4
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Matches 12
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15 US-10-366-709-2
15 US-09-834-765-45
16 US-09-834-765-73
17 US-10-23-187-8
18 US-10-23-187-8
19 US-09-834-765-78
19 US-09-834-765-78
10 US-10-23-187-8
11 US-10-23-187-8
12 US-10-24-652-139-14
13 US-10-024-652-129-15
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15 US-10-024-652-205-16
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; SEQ ID NO 2
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Artificial Sequences
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US-09-730-174A-3
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Best Local S
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 6
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 3
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
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Best Local :
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                Best Local Similarity 91. Matches 11; Conservative
                                                 Query Match
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CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
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TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid;
                                                                                      -09-730-174A-3
                                                                                                                                                                                                                                                       TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid FILE REFERENCE: IMUNE-001A
                                                                                                    OTHER INFORMATION:
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ilarity 100.0%;
Conservative (
                                                                                                    Induces Formation of Antibodies and Isolates
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91.7%;
                                90.2%;
91.7%;
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Pred. No. 0.000
1; Mismatches
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                                Score 55; DB 9;
Pred. No. 0.0017;
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Pred. No.
                  Mismatches
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US-09-730-174A-5
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                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 1
LENGTH: 11
TYPE: PRT
                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 10
                                               Sequence 1, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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   APPLICANT: Schmidt-
APPLICANT: Roth, He
TITLE OF INVENTION:
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Missbichler, Albert
Schmidt-Gayk, Heinrich
Roth, Heinz-Jurgen
NVENTION: Method for Determining Parathormone
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Sequence 5, Application US/09730174A

Patent No. US20020110871A1

GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.

TITLE OF INVENTION: Antibodies and Peptide Antigens for Pro-
TITLE OF INVENTION: Selective Binding Specificity to Bioac
FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
15NOTE: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; FEATURE: ; OTHER INFORMATION: Induces Formation of Antibodies US-09-730-174A-5
                                                                                                                                                  , OTHER INFORMATION: Induces Formation of Antibodies and Isolates US-09-730-174A-1
                                                                                                                                                                                                                                                                                   Sequence 1, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H
FILE REFERENCE: INUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
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                                                                         l Similarity 90.
10; Conservative
VSEIQLMHNLG 11
                                   VSEIQFMHNLG 12
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Pred. No. 0.00
1; Mismatches
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                                                                                           Score 51; I
Pred. No. 0.
                                                                               Mismatches
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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Method for Determining Parathormone
FILE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE OF INVENTION: WIMBER: US/10/166,185
CURRENT APPLICATION NUMBER: US/10/166,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: COT/EP00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION DATE: 1999-12-17
NUMBER OF SEG ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO. 7
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CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
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Best Local Similarity
Thes 9; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
US-10-168-185-7
              Sequence 2, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
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Publication No. US20030175802A1
GENERAL INFORMATION:
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Best Local (
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TYPE: PRT
ORGANISM: Homo sapiens
CURRENT APPLICATION NUMBER: US/10/168,185
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88.9%;
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Pred. No. le+06;
0; Mismatches
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Pred. No. 0.098;
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APPLICANT: Ambruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION WIMBER: POT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local :
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LENGTH: 9
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
TILE REFERENCE: 0609-4570002
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local
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PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR PILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
NUMBER OF SEQ ID NOS: 11
SOPTWARE: FastSEQ for Windows Version 4.0
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TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No. 1e+06;
0; Mismatches
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Pred. No.
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Vannenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
COPPMANDE: DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
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US-10-192-673-7
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US-10-168-185-3
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                                                                                        Sequence 3, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7, Application US/10192673 Publication No. US20030166838A1 GENERAL INFORMATION:
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                       Matches
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1999-10-20 PRIOR APPLICATION NUMBER: U.S. PRIOR FILING DATE: 1998-10-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                     Similarity 66. 6; Conservative
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77.8%;
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Pred. No. 1e+06;
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Pred. No. 1e+06;
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APPLICANT: Juppner, Haraid
APPLICANT: Juppner, Haraid
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Bioactive Peptides and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHr)
TILE OF INVENTION: Hormone-Related Peptide (PTHrP)
TILE REFERENCE: 0609.4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR APPLICATION NUMBER: U.S. 50/105,530
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR PRIORAPPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER: DG SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
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                                                                                                    Sequence 61, Application US/10033741; Publication No. US20030049640A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                             문
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Best Local S
Matches 6
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Best Local
APPLICANT: Herman, et al.
TITLE OF INVENTION: Proteins, Genes and TITLE OF INVENTION: Response
FILE REFERENCE: 9195-079
CURRENT APPLICATION NUMBER: US/10/033,741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gardella, Thomas J. APPLICANT: Kronenberg, Henry APPLICANT: Potts, John T. APPLICANT: Juppner, Harald
                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description of Artificial Sequence: synthetic peptide
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87.5%;
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                                                               Genes and Their Use For Diagnosis and Treatment
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Pred. No. 1e+06;
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Pred. No. 1e+06;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                           DB 14;
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CURRENT FILING DATE: 2001-12-27

NUMBER OF SEQ ID NOS: 80

SOFTWARE: Patentin version 3.0

SEQ ID NO 61

LENGTH: 10

TYPE: PRT

ORGANISM: Homo sapiens

US-10-033-741-61

Query Match
Best Local Similarity 55.6%; Pred. No. 2.1e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Matches 5; Conservative 2; Mismatches 0; Gaps 0;

Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Search completed: May 18, 2004, 10:20:59

Job time: 33.25 secs
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
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Query
Match
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep: *

2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep: *

3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep: *

4: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep: *

5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep: *

6: /cgn2_6/ptodata/2/iaa/backfiles1.pep: *
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                                                                                                                            Copyright
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US-09-442-989-31
US-08-817-547A-1
US-08-817-547A-2
US-09-421-379-6
US-09-421-379-7
US-08-817-547A-4
US-08-817-547A-4
US-08-817-547A-4
US-08-817-547A-4
US-08-817-547A-6
US-08-1482-528-42
US-08-177-109A-5
US-08-177-109A-5
US-08-177-109A-5
US-08-177-109A-5
US-08-177-109A-5
US-08-177-109A-5
US-08-177-547A-1
US-08-177-547A-7
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                                                                                                                            Sequence 31, Appl Sequence 2, Appli Sequence 6, Appli Sequence 3, Appli Sequence 3, Appli Sequence 10, Appli Sequence 10, Appli Sequence 42, Appli Sequence 4, Appli Sequence 4, Appli Sequence 42, Appli Sequence 42, Appli Sequence 56, Appli S
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        Sequence
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US-09-442-989-31
                                                                                                                                                                OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION:
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                                                                                                                                                                                         NAME/KEY: PEPTIDE
LOCATION: (10)
                                                                                                                                                                                                                                                                                                                        NAME/KEY: PEPTIDE
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US-08-428-257A-14	5-579-	15-181-	US-09-556-605-60	US-09-064-750-1	-09-	US-09-091-071-3	US-09-434-122-49	US-09-393-385B-108	0-1	-08	٠,	83-853A-	28B-9	9	16-778E	3-2	83-831B-
Sequence 14, Appl	equence 2,	equence 83,	equence 60	equence 1,	equence 61	equence 3,	equence 49	equence 10	equence 12	equence 12	Sequence 49; Appl	equence 12	equence 97	equence 9,	equence 9,	σ	equence 26, App

ALIGNMENTS

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; FEATURE:
, NAME/KEY: MOD RES
; LOCATION: (8)
; OTHER INFORMATION: N1e
US-09-442-989-31
                                                                                                                                                                                                      OTHELL PEPTIDE NAME/KEY: PEPTIDE LOCATION: (4)
OTHER INFORMATION: G FEATURE:
; NAME/KEY: PEPTIDE
; NAME/KEY: (6)
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LOCATION: (3)
OTHER INFORMATION: Ser(OtBu)
                                                                                                                                     LOCATION: (9)
OTHER INFORMATION: His(Trt)
                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: FMOC-Ala
                                                                                                                                                                                                  Gln(Trt
                                                                                                                                                                                                                                                              Glu (OtBu)
                                                                            Asn (Trt)
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RESULT 2 US-08-817-547A-1

GENERAL INFORMATION:
APPLICANT: Adermann, Knu
APPLICANT: Hock, Dieter

Adermann, Knut

STREET: 191 CONTROL OF STATE: Georgia

CORRESPONDENCE ADDRESS:

ADDRESSEE:

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COUNTK:

ZIP: 30303

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: PATENTIN PC_DOS/MS-DOS
SOFTWARE: PATENTIN DATA;
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA;
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700

104-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 75.4%; Score 46; DB 4; Best Local Similarity 75.0%; Pred. No. 0.013; Matches 9; Conservative 1; Mismatches
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                                                                                                     Sequence 2, Application US/08817547A Patent No. 6030790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                          73.8%; Score 45; DB 3; Length 10; 90.0%; Pred. No. 0.017; Indels
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                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0669.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: unknow MOLECULE TYPE: per HYPOTHETICAL: no ANTI-SENSE: no US-08-817-547A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 8; Conserva
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: peptide
US-09-421-379-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 4
US-09-421-379-6
                                                                                                                         SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/09421379 Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNWER: PCT/EP95/03757
APPLICATION UNWER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGERT T.
REGISTRATION NUMBER: 22,176
REGISTRATION NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE SEARCH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

"PDF:/CATION NUMBER: US/08/817,547A
                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 191 Pea
CITY: Atlanta
STATE: Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 39; DB 3; Pred. No. 3e+05; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
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US-08-817-547A-2

GENERAL INFORMATION:

g Ş

Query Match Best Local S Matches

Similarity 9; Conserv

Conservative

SVSEIQFMHN 10

TOPOLOGY: unknov MOLECULE TYPE: per HYPOTHETICAL: no ANTI-SENSE: no

TYPE: amino acid STRANDEDNESS: un

unknown peptide unknown LENGIH:

10 amino acids

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Sequence 7, Application US/09421379

Patent No. 6495662

GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Fotts, John T.
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bloactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Bloactive Peptides and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO. 7
                                                                                                                                                                                                                                        Sequence 3, Application US/08817547A

Patent No. 6030790
GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Hock, Dieter
APPLICANT: Hock, Dieter
APPLICANT: Hock, Dieter
APPLICANT: Bagerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Se
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Jones & Askew, LLP
STREET: 191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-817-547A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic;
; OTHER INFORMATION: peptide
US-09-421-379-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 5
US-09-421-379-7
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Best Local
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO-postible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A FILING DATE: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                CITY: Atlanta
STATE: Georgi
                                                                                                                                                                       ZIP: 30303
                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 66.7%; 6; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 77.8
7; Conservative
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                                                                                                                                                                                         Georgia
Y: USA
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Pred. No. 3e+05;
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Pred. No. 3e+05;
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US-09-421-379-10
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                                                                                               RESULT 8
US-08-140-137A-42
Sequence 42, Application US/08140137A
Patent No. 5817617
GENERAL INFORMATION:
APPLICANT: TOOMANEN, ELAINE
APPLICANT: MASURE, H. R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Sequence 10, Application US/09421379

Sequence 10, Application US/09421379

Patent No. 6495662

GENERAL INFORMATION:
APPLICANT: Apricantial informas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Kronenberg, Henry
APPLICANT: Juppner, Harald
FITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001

FILE REFERENCE: 0609.4570001

CURRENT APPLICATION NUMBER: US/09/421,379

CURRENT APPLICATION NUMBER: US. 60/105,530

EARLIER APPLICATION NUMBER: US. 500

EARLIER APPLICATION NUMBER: US. 500

EARLIER APPLICATION SAME SAME SAME SAME SAME SEQ ID NO 10

SEQ ID NO 10

LENGTH: 9

TYPE: DEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
TOPOLOGY: unkr
NOLECULE TYPE: I
HYPOTHETICAL: nc
NNTI-SENSE: no
US-08-817-547A-3
                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEFT 1994
ATTORNEY, AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE, DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPAX: 404-818-3700
TELEPAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 3:
                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 SVSEIQFM 8
SVSEHQLLH
                                      SVSEIOFMH
                                                                                47.5%;
nilarity 66.7%;
Conservative
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                                                                                  Score 29; DB 4;
Pred. No. 3e+05;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 31; DB 3;
Pred. No. 3e+05;
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                                                                                       Gaps
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BRUCE E.; WETTENHALL, RICHARD E.H.
TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL
; HYPERCALCEMIA OF MALIGNANCY-PTHRP
NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/715,280
FILING DATE: 14-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 199,235
FILING DATE: 09-MAY-1988
APPLICATION NUMBER:
FILING DATE: 09-MAY-1988
APPLICATION NUMBER:
FILING DATE:
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INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
FENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 5460978
, APPLICANT: MARTIN, THOMAS J.; MOSELEY, JANE M.; KEMP,
                                                                                                                                                                                         SEQ ID NO:3:
                                                                                  Matches
                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC COMPATIBLE
COMPUTER: Floppy disk
COMPUTER: IBM FC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/140,137A
FILLING DATE: 27-MAY-1994
CLASSIFICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 411 ...
STREET: Hackensack
CITY: Hackensack
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
DESCRIPTION: CARD peptide segence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Jackson Egg., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-096
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-5800
TELEPAX: 201 343-1684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                      LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 VSEIQFMH 9
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                                                                                  Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ISEEQFVH 12
                                              SVSEIQFMHN 10
             AVSEHQLEHN 10
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                                                                                    Conservative
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                                                                                                 45.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              49
                                                                                Score 28; DB 6; Length 11; Pred. No. 32; 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 29; DB 2; Length 12; Pred. No. 23; 2; Mismatches 1; Indels
                                                                                    0
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                                                                                    Gaps
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US-08-817-547A-5
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; HYPOTHETICAL: no
; ANTI-SENSE: no
US-08-817-547A-5
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                                                                                                                                                                                                                                                                US-08-817-547A-4
                                                                                                                                                                                                                                                                             RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Jones & Askew, LLP
STREET: 191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 1...
STREET: 1...
CITY: Atlanta
Caporgia
              CITY: Atlanta
STATE: Georgia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
COUNTRY: U
ZIP: 30303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                      1 SVSEIQ 6
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Y: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unknown
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Sequence 5, Application US/08817547A Patent No. 6030790 GENERAL INFORMATION:
Sequence 4, Application US/08817547A
Patent No. 6030790
GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hook, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
PROF APPLICATION DATA:
PRICHATION NUMBER: PCT/EP95/03757
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Adermann, .....APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides
TITLE OF OFFICERS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             404-818-3700
                                                                                                                                                                                                                                                                                                                            100.0%; Pred. ...
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                       42.6%; Score 26;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US/08/817,547A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07826-0007
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Version #1.30
                                                                                                                                                                                                                                                                                                                                                                       DB 3;
3e+05;
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GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: Potts, John T.
APPLICANT: Dotts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: BLOAGATIVE Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT APPLICATION NUMBER: US.
EARLIER APPLICATION NUMBER: US.
EARLIER FILING DATE: 1999-10-20
LOUNDER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NOS: 13
                                                                                                                                                   ; FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic;
OTHER INFORMATION: peptide
US-09-421-379-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 12
US-09-421-379-8
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APPLICATION NUMBER: PCT/EP95/(
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782(
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3709
TELEPHONE: 404-818-3799
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                      Query Match
Best Local Similarity
Matches 5; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8, Application US/09421379 Patent No. 6495662
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRICE APPLICATION HOMBER: US/08/817,547A
                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL:
ANTI-SENSE: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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STRANDEDNESS: unl
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                                 SVSEIQFMH 9
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100.0%; Pred. No.
                                                                                            42.6%;
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                                                                                          Score 26; DB 4;
Pred. No. 3e+05;
                                                                           Mismatches
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US-08-482-228-42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Tseng-Law, Janet
APPLICANT: Kobori, Joan A.
APPLICANT: Al-Abdaly, Fahad A.
APPLICANT: Guillermo, Roy
                                                  APPLICANT: Guillermo, Roy
APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
NUMBER OF SEQUENCES: 215
CORRESPONDENCE ADDRESS:
ADDRESSEE: Janice Guthrie, Ph.D.
STREET: P.O. Box 15210
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Patent No. 5900 ...
Parent No. 5900 ...
PREPARENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (714) 440-5353
TELEPAX: (714) 553-1952
INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #10, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,228
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 42, Application US/08482228 Patent No. 5968753
Sequence 42, Application US/08482528 Patent No. 6017719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Tseng-Law, Janet
APPLICANT: Kobori, Joan A.
APPLICANT: Kobori, Joan A.
APPLICANT: Al-Abdaly, Fahad A.
APPLICANT: Guillermo, Roy
APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
NUMBER OF SEQUENCES: 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Guthrie, Janice
REGISTRATION NUMBER: 35,170
REFERENCE/DOCKET NUMBER: IT-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (714) 440.5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Janice Guthrie, Ph.D.
STREET: P.O. Box 15210
                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 92713-5210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: P.O. Box CITY: Irvine
STATE: California
                                                                                                                                                                                                                                                  / Match 41.0%;
Local Similarity 57.1%;
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                                                                                                                                                                                  3 SEIQFMH 9
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                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                               linear
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                                                                                                                                                                                                                               1; Mismatches
                                                                                                                                                                                                                                                  Score 25; DB 2;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                       Length 12;
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COUNTRY:

CITY: Irvine STATE: California

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COMPUTER: USA

ZIP: 92713-5210

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.25

SOFTWARE: PACENTIN RELEASE #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,528
FILING DATE: 07-UM-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Guthrie, Janice
REGISTRATION NUMBER: IT-4630CIP4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (714) 440-5553
TELEPAX: (714) 453-1952
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 15
US-08-177-109A-56
(Sequence 56, Application US/08177109A
; Patent No. 5869615
; Patent No. 5869615
; Patent OF SEQUENCE BORNES B. Hourcade and Teresa J. Oglesby
TITLE OF INVENTION: MODIFIED COMPLEMENT PROTEASES
; NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ß
CORRESPONDENCE ADDRESSEE: PATERA L. PabSt

ADDRESSEE: Patera L. PabSt

STREET: 2800 One Atlantic Center

STREET: 1201 West Peachtree Street

CITY: Atlanta

STATE: Georgia

COUNTRY: USA

ZIP: 30309-3450

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compatible

OPERATION SYSTEM: PC-DOS/MS-DOS

SOCTWARRE: PALEITION PATA:

APPLICATION NUMBER: US/08/177,109A

FILING DATE: 03-JAN-1994

CLASSIFICATION NUMBER: US/08/177,109A

FILING DATE: 03-JAN-1994

ATTORNEY/AGENT INFORMATION:

NAME: PabSt, Patrea L.

REGISTRATION NUMBER: 31,284

REFERENCE/DOCKET NUTWER: WU 107

TELECOMMUNICATION INFORMATION:

TELEPHONE: 4040 873-8794

TELEPAN: (404) 873-8795

INFORMATION FOR SEQ ID NO: 56:

SEQUENCE CHARACTERISTICS:

LENGTH: 5 amino acids

TYPE: amino acids
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Best Local Similarity 57.1%;
Matches 4; Conservative 1
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Search completed: May 18, 2004, 10:06:44
Job time : 13.75 secs
                                                                                                                                                                                                                        MOLECULE TYPE: protein; HYPOTHETICAL: NO US-08-177-109A-56
                                                                           Вb
                                                                                                                                            Query Match 39.3%; Score 24; DB 2; Length 5; Best Local Similarity 60.0%; Pred. No. 3e+05; Matches 3; Conservative 2; Mismatches 0; Indels
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Result
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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score
and is
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Gapop 10.0 , Gapext 0.5
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112.614 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query
Match
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59
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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                                                                          Copyright
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GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd
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S29735
PW0002
A56029
E49033
A38841
B41835
A40795
I38888
I38888
I38888
FP0231
PT0231
PT0231
SPPGNK
S28055
S278055
S278053
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A37268
PH1190
PH1187
PH1189
A47364
PH0924
S25266
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B46453
A55837
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NADH2 dehydrogenas
Ig heavy chain CRD
neuromedin K - pig
cytochrome b559 co
hypothetical prote
ranatachykinin C -
                                                                                                                                   translation elonga
glycoprotein H-a -
COI intron 16 prot
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	111 112 112 112 113 113 113 113 113 113
	<i>αα</i> 4 αααααααααααα
ALIGNMENTS	\$60354 \$42900 \$42909 \$25485 \$71034 \$71034 \$71175 \$71035 \$1174 \$11173 \$114206 \$114306 \$
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	retinal oxidase - acidic proline-ric hypotherical prote transcription fact transcription fact complex polypept potB protein - Sal T-cell receptor al T-cell receptor al glutamine-tRNA lig angiotensin-conver hypothetical prote leucokinin VI - Ma IG H chain V-D-J r coat protein beta sperm-activating Ig heavy chain CDR

RESULT 1 S21205 Ig heavy chain V region - human C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Accession: S21205 R;Makiya, R.; Stigbrand, T. Bur. J. Biochem. 205, 341-345, 1992 A;Title: Placental alkaline phosphatase has a binding site for the human immunoglobulin-A;Reference number: S21205; MUID:92209522; PMID:1555592 A;Accession: S21205 Query Match Best Local S Matches 4 1 EVOLVESGG 9 4; Similarity EIQLMHNLG 12 Conservative 35.6%; Score 21; DB 2; Length 12 Pred. No. 6.9e+02; 3; Mismatches 2; Indels Length 12, 0 Gaps 0

Qy 3 SEIQ 6 Db 2 SEIQ 5 RESULT 3 \$78420	Query Match 30.5%; Score 18; DB 2; Length 7; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	RESULT 2 A58718 C;Species: Carnobacterium sp. (fragment) C;Species: Carnobacterium sp. C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998 C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998 C;Accession: A58718 R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, I.F. A;File: Purification and characterization of a new bacteriocin isolated from a Carnobac, File: number: A58718; MUID:92321768; PMID:1622206 A;Reference number: A58718 A;Accession: A58718 A;Realdues: 1-7 <sto> C;Keywords: antibiotic; lanthionine</sto>
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ribosomal protein RL41, mitochondrial [validated] - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000 C;Accession: S78420 R;Goldschmidt-Reisin, S;; Graack, H.R.
                                                                                                                                                                                                                A; Variety: subtype adr C; Date: 18-Um-1993 #sequence_revision 08-Nov-1996 #text_change 15-Aug-1997 C; Accession: B46453 R; Takahashi, K.; Kishimoto, S.; Ohori, K.; Yoshizawa, H.; Machida, A.; Ohnur J. Immunol. 147, 3156-3160, 1991 A; Title: Molecular heterogeneity of e antigen polypeptides in sera from car: A; Reference number: A46453; MUID:92013147; PMID:1717588 A; Accession: B46453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, M.; Yamagud Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and its related peotus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Reference number: A60527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sperm-activating peptide (Tyr-2, Asn-3, Gly-5, Ile-9, Asp-10 SAP-I) - slate-pencil
C;Species: Heterocentrotus mamillatus
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
C;Accession: A60589
                                                                                                                                                                                                                                                                                                                                                                                          e antigen p20e precursor - hepatitis B virus (subtype adr) (fragment)
N;Alternate names: HBe antigen precursor
N;Contains: e antigen
C;Species: hepatitis B virus, HBV
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A;Residues: 1-10 <YOS>
C;Superfamily: unassigned animal
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A;Note: the protein is designated as mitochondrial ribosomal C;Keywords: mitochondrion; protein biosynthesis; ribosome
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A; Accession: S78420
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                                                                                     Experimental source: subtype adr;
Note: sequence extracted from NCBI backbone (NCBIP:60243);
1-10/Domain: signal sequence (fragment) #status predicted;
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Pred. No. 2.1e+03;
                       Score 18; DB 2;
Pred. No. 2.1e+03;
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A37268
A37268
Ig heavy chain C region (129) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Accession: A37268
C;Accession: A37268
C;Accession: A37268
A;Title: Heavy and light chain variable region sequences and ant A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: A37668
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C;Bate: 19-Gan-1996 #sequence_revision 19-Jan-1996 #text_change 17-Mar-1999
C;Accession: A55837
R;Firestine, S.M.; Davisson, V.J.
Blochemistry 33, 11917-11926, 1994
A;Title: Carboxylases in de Novo purine biosynthesis. Characterization of the Gallus gal A;Reference number: A55837; MUID:95001903; PMID:7918410
A;Accession: A55837
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A55837
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997
C;Accession: I46868
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A;Molecule type: protein
A;Residues: 1-12 <FIR>
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                                                                     A; Molecule type: mRNA
A; Residues: 1-10 < RUF >
                                                                                       A;Status: preliminary A;Molecule type: mRNA
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A; Residues: 1-7 < FRI>
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T-cell receptor alpha chain V region (Cw3/IFI1) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Jate: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1187
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell A;Reference number: $26512; MUID:92364546; PMID:1380061
A;Accession: PH1187
A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
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A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1190
A;Accession: PH1190
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C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1189
C;Accession: PH1189
C;Accession: J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; B:
J; Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T (A;Reference number: S26512; MUID:92364546; PMID:1380061
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A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
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RESULT 12
A47364
A47364
placental lactogen-I precursor - mouse (fragment)
C;Species: Mus sp. (mouse)
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996
C;Accession: A47364
R;Shida, M.M.; Ng, Y.K.; Soares, M.J.; Linzer, D.I.
Mol. Endocrinol. 7, 181-188, 1993
A;Title: Trophoblast-specific transcription from the mouse placental lactog A;Title: Trophoblast-specific transcription from the mouse placental lactog A;Reference number: A47364; MUID:93225959; PMID:8469232
A;Accession: A47364
A;Status: preliminary; translated from GB/EMBL/DDBJ
A:Malacula type: DNA
                                                                                                                                         pilE protein - Escherichia coli (fragment)
(/Species: Escherichia coli
(/Species: Escherichia coli
(/Species: Escherichia coli
(/Species: Escherichia coli
(/Date: 28-May-1993 #sequence_revision 28-May-1993 #text_change 11-Jun-1993
(/Accession: S25266
R;Dupuy, B.; Taha, M.K.; Possot, O.; Marchal, C.; Pugsley, A.P.
Mol. Microbiol. 6, 1807-1894, 1992
Mol. Microbiol. 6, 1807-1894, 1992
A;Title: PulO, a component of the pullulanase secretion pathway of Klebsiella oxytoca,
A;Reference number: S25266; MUID:92374839; PMID:1354833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A, Experimental source: concanavalin A-activated lymphoblast C_1 Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.

J. Exp. Med. 174, 1467-1476, 1991

A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental allergy

A;Reference number: PH0891; MUID:92078857; PMID:1836012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Accession: PH0924
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
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A; Residues: 1-10 < RES>
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A; Residues: 1-11 <GOL>
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                                                                   C; Genetics:
                                                                                       A;Molecule type: protein
A;Residues: 1-7 <DUP>
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Pred. No. 5.1e+0
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 12
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Perfect score:
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         332226543210
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Gapop 10.0 , Gapext 0.5
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59
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ALLAC CASUP
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CD11 LITXA
COMB PETAM
RS19 TOBB
LCKA LEUMA
ANOSH CLYJA
PPEA
MOSH CLYJA
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ALIGNMENTS

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SEQUENCE. SEQUENCE. TISSUE=Larva; TISSUE=Larva; TISSUE=Arva; TISSUE=Arva; Thouse 4.; Thouse 5.; Thouse 5.; Thouse 5.; Thouse 6.; Tho	_CYDPO STANDARD; PRT; 8 AA. ALL6 CYDPO STANDARD; PRT; 8 AA. P82157; 30-MAY-2000 (Rel. 39, Created) 30-MAY-2000 (Rel. 39, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update) Cydiastatin 6. Cydiastatin 6. Cydiastatin 6. Cydia pomonella (Codling moth). Cydias pomonella (Codling moth). Cydiastatin 5. Cydiastatin 6. Cydiast	Query Match 30.5%; Score 18; DB 1; Length 7; Best Local Similarity 100.0%; Pred. No. 1.4e+05; Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 3 SEIQ 6 6 2 SEIQ 5	ARUIT ANC CARUI STANDARD; PRT; 7 AA. 36960; 1-JUN-1994 (Rel. 29, Created) 1-JUN-1994 (Rel. 29, Last sequence update) 6-OCT-2001 (Rel. 40, Last annotation update) antibiotic carnocin UI49 (Fragment). armobacterium sp. (strain UI49). acteria; Firmicutes; Lactobacillales; Carnobacteriace armobacterium. CBI TaxID=35782; EQUENCE. EDLINE=93321768; PubMed=1622206; EQUENCE SIF.; Carnobacterium sp."; Carnobacterium sp."; pli Environ and characterization of a new bacterioci Carnobacterium sp."; ppl. Environ. Microbiol. 58:1417-1422(1992). 1 - FUNCTION: Lanthionine-containing peptide antibioti 1 - FUNCTION: Lanthionine-containing intibiotic; Bacteriocin; Lantibiotic. ON TER 7 27 20UENCE 7 AA; 786 MW; 741776D05B05B810 CRC64;

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RESULT 3
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MOD RES
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15-JUL-1998 (Rel. 36, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
10-OCT-2003 (Rel. 42, East annotation update)
Allergen Fus s I3596* (Fragment).
Fusarium solani (subsp. plsi) (Nectria haematococca).
Eukaryota, Fungi, Ascomycota; Pezizomycotina; Sordariomycetes;
Hypocreomycetidae; Hypocreales; Nectriaceae; Nectria.
NCBI TaxID=70791;
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NON TER
SEQUENCE
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Verma J., Gangal S.V
Submitted (JUL-1997)
                                                                                                                              MEDLINE=98291870; PubMed=9629918; Flengsrud R., Skjeldal L.; "Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5."; Electrophoresis 19:802-806(1998).
-!- FUNCTION: This protein promotes the GTP-dependent translocation of the CTP-dependent translocation.
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InterPro;
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                     ribosome.
SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the GTP-binding
EF-G/EF-2 subfamily.
                                                                                                         the nascent protein chain from
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. 36, Last
. 41, Last
r G (EF-G)
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S.V.;
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40.0%;
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an allergic re
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Last sequence
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0B2879C45B573767 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           annotation update) (CP 5) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C372C441F5B69041 CRC64;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16; DB 1;
Pred. No. 1.4e+05;
2; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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1.4e+05;
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A-site to the N
                                                                                                                A-site
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                                            elongation factor family
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P-site of the
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    RESULT 6
TRPI PSEPU
ID TRPI PS
AC P36414
AC P36414
AC O1-JUN
DT 01-JUN
DT 01-JUN
DT 15-MAR
DE HTH-ty,
DE actival
GN Pseudo
OC Bacter
OC Pseudo
OC Bacter
OC Pseudo
OC STRAIN
RP SEQUEN
RP SEQUEN
RP SEQUEN
RA EBERTY
RT PUNA s
RT PUNA s
RT PULIAG
RT Biochi
CC -!- FU
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PVK1_PE
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Best Local S
Matches 3
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Best Local
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Elongation factor; Protein biosynthesis;
NON_TER 11 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOD RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Periplaneta americana (American cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; E
Blattidae; Periplaneta.
NCBI TaxID=6978;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PVK1 PERAM STANDARD; PRT; 11 AA P41837; P41837; O1-NOV-1995 (Rel. 32, Created) O1-NOV-1995 (Rel. 32, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update periviscerokinin-1 (Pea-PVK-1).
                                                                                                                                                                                                                                                                                                                                      01-UUN-1994 (Rel. 29, Created)
01-UUN-1994 (Rel. 29, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
HTH-type transcriptional regulator trpI (TrpBA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Abdominal perisympathetic organs;

MEDLINE=9523021; PubMed=7716075;

Predel R., Linde D., Rapus J., Vettermann S., Penzli
"Periviscerokinin (Pea-PVK); a novel myotropic neuro
"Periviscerokinin of the American cockroach.";

Perivides 16:61-66(1995).

i- FUNCTION: MYOACTIVE PEPTIDE; HAS EXCITORY ACTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PERAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                  Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales
Pseudomonadaceae; Pseudomonas.
NCBI_TaxID=303;
                                                                                                                                                                                                                                                                                                                                                                                                        P36414;
01-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neuropeptide; Amidation.
         Biochimie 71:521-531(1989)
                                                   putida.
                                                                        Eberly L., Crawford I
                                                                                                                   MEDLINE=89335826; PubMed=2503057;
                                                                                                                                          SEQUENCE FROM N.A. STRAIN=PPG1 C1S;
                                                                                                                                                                                                                                                                           Pseudomonas putida.
                                                                                                                                                                                                                                                                                                                   activator)
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3; Conserv
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25.0%;
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                                                                      tryptophan
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Pred. No. 3.5e+03;
3; Mismatches 3
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           EXPRESSION
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Pred. No. 5.5e+03
1; Mismatches
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39DB5419D7605728
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                                                                           synthase
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              TRPBA
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Blattoidea;
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                GENES ENCODING
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AL18_CAL18_C
AC P81821
DT 30-MAY
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DT 30-MAY
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Best Local S
Matches 2
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no was modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@isb-sib.ch).
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30-MAY-2000
30-MAY-2000
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InterPro; IPR000847; HTH_LYSR.
PROSITE; PS50931; HTH_LYSR; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUB-Larva;
TISSUB-Larva;
MEDLINE-98054539; PubMed-9392829;
Dave H., Johnsen A.H., Maestro J.-L., Scott
Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin peptides 18:1301-1309(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ALL7 CYI
P82158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cydia pomonella (Codling moth).
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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                                                                                                                                                                                                                                                                                                                                                                                                                       "I- SIMILARITY: Belongs to the allatostatin Neuropeptide; Amidation.
MOD_RES 7 amrnamrow
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                                                                 P81821;
30-MAY-2000
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    Carcinustatin
                       30-MAY-2000
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SIMILARITY: Contains 1 HTH lysR-type DNA-binding domain.
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                                                                                                               CARMA
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                                                                                                                                                                                                                                                                                                                                   Similarity
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P82152;
30-MAY-2000
30-MAY-2000
30-MAY-2000
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                            P83281;
28-FEB-2003 (Rel
28-FEB-2003 (Rel
28-FEB-2003 (Rel
FMRFamide-like r
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Cerebral ganglion, and Thoracic MEDLINE=98121193; PubMed=9461295; Duve H., Johnsen A.H., Maestro J.-L., S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carcinus maenas (Common shore crab) (Green crab).
Eukaryota, Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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MOD RES 8 8
SEQUENCE 8 AA; 934 MW
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-i- SIMILARITY: Belongs to the allatostatin
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TISSUE=98054539; PubMed=9392829;
MEDLINE=98054539; PubMed=9392829;
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(Rel. 41,
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(Rel. 39, Last sequence update)
(Rel. 39, Last annotation update)
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                            el. 41, Last sequence update)
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neuropeptide FLP8 (VSHNNFLRF-amide)
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2; Mismatches
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Pred. No. 1.4e+05
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Macrobrachium rosenbergii

(Giant

fresh water prawn)

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Gaps

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RESULT 12
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AC PA185
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DT 30-MF
DT 30-MF
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UF02 MOUSE
P38640;
                                                -AL19 CARMA
P81822;
P81842-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-21107394; PubMed=11179812; Sithigorngul P., Saraithongkum W., Longyant S., Panchan Sithigorngul W., Petsom A.; "Three more novel FMRFamide-like neuropeptide sequences eyestalk of the giant freshwater prawn Macrobrachium ros peptides 22:191-197(2001).

-i- MASS SPECTROMETRY: MW=1133.8; METHOD=MALDI.
-i- SIMILARITY: Belongs to the FARP (FMRFamide related programme programme).
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Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Caridea
Palaemonoidea; Palaemonidae; Macrobrachium.
NCBI TaxID=79674;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
15-VAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of fibroblasts (P32) (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neuropeptide; Amidation.
MOD_RES 9 9
SEQUENCE 9 AA; 1133 M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (Mouse).
 Carcinus maenas
                  Carcinustatin 19.
                                     30-MAY-2000
                                                                                                                                                                                                                                                                                                                                     SEQÜENCE
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X MEDLINE=$3099270; PubMed=1463853;

X KUNITZ M., Camara B., Weil J.-H., Schantz R.;

A KUNITZ M., Camara B., Weil J.-H., Schantz R.;

T "The psbi gene from bell pepper (Capsicum annuum): plastid R!

T editing also occurs in non-photosynthetic chromoplasts.";

L plant Mol. Biol. 20:1185-1188 (1992).

C -!- FUNCTION: This b-type cytochrome is tightly associated w
reaction center of photosystem II and possibly is part o
water-oxidation complex.

C water-oxidation omplex.

C :-- SUBUNIT: Heterodimer of an alpha subunit and a beta subun

C :-- SUBUNITY: Belongs to the psbE / psbF family.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PSBF_CAPAN
Q03367;
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NCBI_TaxID=6759;
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Eubrachyura; Portunoidea; Portunidae; Carcinus.
       PROSITE; PSOC
Chloroplast;
NON_TER
                Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
lamiids; Solanales; Solanaceae; Capsicum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last Sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Cytochrome b559 beta subunit (PSII reaction c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- FUNCTION: May act as a neurotransmitter or neu-
-!- SIMILARITY: Belongs to the allacostatin family
Neuropeptide, Amidation, Multigene family
MOD_RES 10 10 AMIDATION (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Isolation and identification of multiple neuropeptides allatostatin superfamily in the shore crab Carcinus maen Eur. J. Biochem. 250:727-734(1997)
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Chloroplast.
Eukaryota; Viridiplantae;
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                                                                                                                                               use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and foentities requires a license agreement (See http://www.isb-siborsend an email to license@isb-sib.ch).
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                                                                                              EMBL; X65570; CAA46539.1;
PIR; S28055; S28055.
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                                                                                                                                                                                                 s SWISS-PROT entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EW European Bioinformatics Institute. There are no restr by non-profit institutions as long as its content
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         ; PS00537; CYTUCHANIAST; Photosystem 1
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2; Conserv
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30-MAY-2000
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10-OCT-2003
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DOMAIN
SEQUENCE
TISSUE=Intestine;
MEDLINE=91254337; PubMed=2043143;
MEDLINE=91254337; PubMed=2043143;
Kozawa H., Hino J., Minamino N., Kangawa K., Matsuo H.;
"Isolation of four novel tachykinins from frog (Rana ca brain and intestine.";
Biochem. Biophys. Res. Commun. 177:588-595(1991).
                                                                                                                                                                                                                                                                                  TKNC_RANCA STANDARD; PRT; 10 AA. P22690; PC 1991 (Rel. 19, Created) 01-AUG-1991 (Rel. 19, Last sequence update) 10-CCT-2003 (Rel. 42, Last annotation update) Ranatachykinin C (RTK C).
Rana catesbeiana (Bull frog).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Skin secretion;
MEDLINE=97175050; PubMed=9022710;
Simmaco M., Mignogna G., Canofeni S.,
                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana
NCBI_TaxID=8400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the brevinin family.
Amphibian defense peptide; Antibiotic; Amidation.
MOD_RES 10 10 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rana temporaria (European common frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
NCBI_TaxID=8407;
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temporaria."
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                                                                                                                                                                  SEQUENCE, AND SYNTHESIS.
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Pred. No. 7.7e+03;
4; Mismatches 2;
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LUMENAL (POTENTIAL).
817D0F59D6D69DC5 CRC64;
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Pred. No. 7.7e+03;
1; Mismatches 1;
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Search completed: May 18, Job time: 6.75 secs
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TISSUE=Intestine;
MEDLINE=94023216; PubMed=8210506;
MEDLINE=94023216; PubMed=8210506;
Kangawa K., Kozawa H., Hino J., Minamino N., Matsuo H.;
Kangawa K., Kozawa H., Hino J., Minamino N., Matsuo H.;
"Four novel tachykinins in frog (Rana catesbeiana) brain and
intestine.";
Regul. Pept. 46:81-88(1993).
-!- FUNCTION: Tachykinins are active peptides which excite neurons,
evoke behavioral responses, are potent vasodilators and
secretagogues, and contract (directly or indirectly) many smoot)
                                                                                                                                                                                                                    PIR; C61033; C61033.
InterPro; IPR002040; Tachy Neurokinin.
PROSITE; PS00267; TACHYKININ; 1.
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                                                                                                                                                                                        Tachykinin; Neuropeptide; Amidation.
MOD_RES 10 10 AMIDATI
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-!- SIMILARITY: Belongs to the tachykinin family.
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Pred. No.
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           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ALIGNMENTS

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Q9BS71 PRELIMINARY; PRT; 11 AA.
Q9BS71
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2000 (TrEMBLrel. 21, Last annotation update)
Pibrinolytic metalloproteinase (Fragment)
Agkistrodon contortrix.
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Agkistrodon.
NCBI TaxID=8720;
                                                                                                                                                                                                                                                                                                                                     of the southern copperhead snake (Agkistrodon contortrix)."; contortrix). Biophys. 289:197-207(1991). NON TER 11 11 SEQUENCE 11 AA; 1209 MW; 7CA02D1D41E8772B
                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE.
SEQUENCE.
MEDILINE=91378546; PubMed=1898066;
Guan A.L., Retrios A.D., Henderson G.N., Markland F.S.Jr.;
Guan A.L., and characterization of a fibrinolytic enzyme
"Purification and characterization of a fibrinolytic enzyme
                                                                                                                                                                                                                                                    Similarity
5; Conserve
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71.4%;
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Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
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-i- MASS SPECTROMETRY: MWH=1236; METHOD=FAB.
SEQUENCE 12 AA; 1236 MW; 147AA70FDF472724 CRC64;
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Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structure of the dynastins from
"Peptides from Australian frogs. The structure of the dynastins from
Limnodynastes salmini and Fletcherin from Limnodynastes fletcheri.";
Aust. J. Chem. 46:1235-1244(1993)
-i- MASS SPECTROMETRY: MW=7772; METHOD=FAB.
SEQUENCE 8 AA; 772 MW; 7B58772455A2C728 CRC64;
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Raftery M.J., Bradford A.M.,
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DYNASTIN 3.
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                                                                     Limnodynastes salmini (Salmin's-striped fr
Eukaryota; Metazoa; Chordata; Craniata; Ve
Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
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NCBI_TaxID=39404;
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Neobatrachia; Bufonoidea; Myobatrachi
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2; Mismatches
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Pred. No. 1e+06;
1; Mismatches
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DYNASTIN 7.
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Q34909 PRELIMINARY;
Q34909;
Q1-NOV-1996 (TrEMBLrel. 01, C
01-NOV-1996 (TrEMBLrel. 01, I
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Cytochrome b (Fragment).
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Bradford A.M., Raftery M.J., Bowle J.H., The Structure of the Gy
"Peptides from Australian frogs. The structure of the Gy
"Inmodynastes salmini and Fletcherin from Limnodynastes
Limnodynastes salmini and Fletcherin from Limnodynastes
Aust. J. Chem. 46:1235-1244(1993).

Aust. J. Chem. 46:1235-1244(1993).

Aust. J. Chem. 46:1235-1248(1993).

Aust. J. Chem. 46:1235-1248(1993).

Aust. J. Chem. 46:1235-1248(1993).

Aust. J. Chem. 46:1235-1248(1993).
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TISSUE=SKIN SECRETION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.; "Peptides from Australian frogs. The structure of the dynastins from Limnodynastes salmini and Fletcherin from Limnodynastes fletcheri."; Aust. J. Chem. 46:1235-1244(1993).

-I- MASS SPECTROMETRY: MW=1114; METHOD=FAB.
SEQUENCE 12 AA; 1114 MW; 3AB5A976CAA72728 CRC64;
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TISSUE-SKIN SECRETION;
Bradford A.M., Raftery M.J., Bow
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Ev
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea;
Limnodynastinae; Limnodynastes.
SEQUENCE FROM N.A.

MEDLINE=8823478; PubMed=2836084;

McCracken A., Uhlenbusch I., Gellissen G.;

"Structure of the cloned Locusta migratoria mitochondrial

"Struction mapping and sequence of its ND-1(URF-1) gene.

Curr. Genet. 11:625-630(1987).

EMBL; X05286; CAA28905.1; -.

GO; GO:0005739; C:mitochondrion; IEA.
                                                                                                                                                                                        Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha; Acridoidea; Acrididae; Oedipodinae; Locusta.
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Vertebrata; Euteleostomi;
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Liermann H., Harder T., Haas L.;

Liermann H., Harder T., Haas L.;

I figenetic analysis of the central untranslated genome reg

Proximal coding part of the F gene of wild-type and vacc

T morbilliviruses.";

RL Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases

REMBL; AF026237; AAC09167.1; -.

DR EMBL; AF026237; AAC09167.1; -.

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01-AUG-1998 (TrEMBLrel. 07, C:
01-AUG-1998 (TrEMBLrel. 07, L:
01-DEC-2001 (TrEMBLrel. 19, L:
Fusion protein (Fragment).
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STRAIN-Dog #10757/96;

Chermann H., Harder T., Haas L.;

"Genetic analysis of the central untranslated genome region

"Genetic analysis of the F gene of wild-type and vaccine

morbilliviruses."

Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.

EMBL, AF026234; AACO9164.1; -.
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
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01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
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NCBI_TaxID=11232;
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01-NOV-1996
01-OCT-2002
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Q9PXC3;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
E antigen P20E (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI TaxID=10407;
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Oryza sativa (Rice).
Oryza, Embryophyta; Mrantoideae; Oryza.
Ehrhartoideae; Oryzeae; Oryza.
OCBI_TaxID=4530;
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                                                                                                                                                                                                                                                                            MEDIINE=92013147; PubMed=1717588; Takahashi K., Kishimoto S., Ohori K., Yoshizawa H., Ma Ohnuma H., Tsuda F., Munekata E., Miyakawa Y., Mayumi "Molecular heterogeneity of e antigen polypeptides in carriers of hepatitis B virus.";
J. Immunol. 147:3156-3160(1991).
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MEDLINE=91078641;
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Avian carcinoma virus.
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"3'-terminal region of avian carcinoma virus MH2 shares sequence elements with avian sarcoma viruses Y73 and SR-A.";
J. Virol. 52:703-705(1984).
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C9GD12;
PREMALE. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Ribosomal protein S16 (Fragment).
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01-NOV-1996
01-NOV-1996
01-DEC-2001
Linospadix monostachya.
Chloroplast.
Eukaryota, Viridiplantae, Streptophyta, Embryophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Arecaceae,
                                                                                                                                                                                                                                                                                                                                                                                      "Isolation of chromosome-specific genes arrayed cDNAs and cosmid libraries."; Hum. Mol. Genet. 0.0-0(1995).
EMBL: L32131; AAA73881.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lee C.-C., Yazdani A., Wehnert
Coolbaugh M.I., Chinault C.A.,
Caskey C.T.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; MCBI_TaxID=9606;
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Homo sapiens (Human)
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MEDLINE=84221901; PubMed=6328491;
Friedman D.J., Umeda P.K., Sinha
Rabinowitz M.;
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Mammalia; Eutheria;
NCBI_TaxID=9986;
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(TrEMBLrel. 01, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
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eavy chains.":
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U.S.A. 81:3044-3048(1984).
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Baldini A.,
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GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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Aab21270 Fertilin	Aar64520 RF-2 pept	Aar89301 Japanese	Abu54029 Human DNA	98581 F	98582 F	98580 F prot	Aab86222 Human par	Aar91647 Human par	Aab01866 PTH(1-14)		Aab07467 Antigenic	Aar91646 Human par	Aay97062 PTH-rP N-	Aay78849 Parathyro	01864 PTH	Abp71482 Parathyro	b96891 Rat	71483 Pa	Abp71485 Parathyro

ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hormone N-terminal signaling domain
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New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass. for

(GARD/) GARDELLA T J. (KRON/) KRONENBERG H M. (POTT/) POTTS J T. (JUEP/) JUEPPNER H.

) KRONENBERG H M.) POTTS J T.) JUEPPNER H.

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WPI; 2000-452384/39.

Gardella TJ,

Kronenberg

MH,

Potts JT,

Jueppner

Claim 4; Page 92; 119pp; English.

Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH), L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R_1 is the PTH-1 receptor signal sequence, and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP

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RESULT
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Best Local
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72607
The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, being formed from the N-terminus of parathyroid hormone (PTH). Also included are; (1) a method for producing antibodies useful in the determination PTH levels in a biological sample comprising; (a) providing at least on first peptide antigen comprising a peptide fragment of PTH; (b) administering the first peptide antigen to a host animal to induce antibody production; (c) monitoring the antibody titre produced; (d) isolating antisera produced in the host animal; and (e) selecting antisera from the isolated antisera produced in the host that is capable of binding to a second peptide antigen; (2) an antibody (ab) produced be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11
                                                                                                                                                                                                 New
and
                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in a mammalian cell having PTH-1 receptors, or screening for a peptide non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH nor PTHrP which avoids the need integrated injections to treat osteoporosis
                                                                                                                                                 Claim 1;
                                                                                                                                                                                    determining
                                                                                                                                                                                                                                                            Zahradnik RJ,
                                                                                                                                                                                                                                                                                                                        05-DEC-2000;
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Misc-difference
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                                                                                                                                                                                                                                                                                                 (ZAHR/)
                                                                                                                                                                                   parathyroid hormone (PTH) antigenic peptide inducing the isolation of antibodies having an affinity to it, useful ermining bioactive PTH levels in serum, plasma and/or cell
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LAVIGNE J R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hormone
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                                                                                                                                                 5; llpp;
                                                                                                                                                                                                                                                             Lavigne
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100.0%; Pr
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                    Parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG72608 standard; peptide; 12
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                                                                           WPI;
                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                     primary
                                                                                                                                                                                                                                                                                                                                                                                              Human; parathyroid hormone;
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                                                                                                                                                                                                                                                                                                            Canis
                                                                                                Zahradnik
                                                                                                                                                    05-DEC-2000;
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                                                                                                                                                                                                15-AUG-2002.
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                                                                                                                                                                                                                                                                                                                                                                  Homo
                                                                                                                                (ZAHR/)
                                                                                                                                                                                                                                                                                                                              taurus.
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                                                                                                                                                                                                                                                                                                                                                                                    hyperparathyroidism;
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                                                                                                                    ZAHRADNIK R J.
LAVIGNE J R.
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                                                                                                                                                     2000US-00730174.
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                                                                                               Lavigne
                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                          /label= Leu,
                                                                                                                                                                                                                                                               label= Ser, Ala
                                                                                                                                                                                                                                                                                                                                                                                                                     antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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; mouse; r
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                     peptide 1-12
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                                                                                                                                                                                                                                                                                                                                                                                     rat;
                                                                                                                                                                                                                                                                                                                                                                                              hypercalcaemia;
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                                                                                                                                                                                                                                                                                                                                                                                    bovine; porcine;
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Claim 2;

Page

5; llpp;

English

media.

New parathyroid hormone (PTH) antigenic peptide inducing the and isolation of antibodies having an affinity to it, useful determining bloactive PTH levels in serum, plasma and/or cell

for formation culture

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ARRESULT 4
ANAR9164
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Best Local S
Matches 10
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                                                                                generated using them forms of the hormone
                                                                                                           New antigenic peptide(s) from generated using them, able to
                             Claim
                                                                                                                                                                                                                                                                                                                                                                    28-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
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                             5pp; German
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90.9%;
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Pred. No.
                                                                                                           human parathyroid hormone - and distinguish between active and
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                                                                                                                                      parathyroid hormone - and antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10 AA;
                                                                                                                                                                                                                                      peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin
                                                                                                                                                                                                                                                                                                                 The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipophilicity. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 8; 18pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New parathyroid hormone fragment peptides, in topically applied cosmetic compositions excessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-171243/15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amino acids 1-10 of a parathyroid hormone (pTH)
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                                                                                                                                                                 Sequence
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          used as lipolysis stimulants for slimming treatment of
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                                                                                                                                                                                                                                                                                 This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that creognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or content of the content of cunlike known assays) recognizes that the comment of (A) shorter than the complete (84 aa) peptide are continued and (ii) that apparently intact peptide may be biologically content of the content of complete (A) and also takes into account the fact that some fragments of (A) care antisyonistic (these have the receptor-binding site but lack the N-contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                               Query Match
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Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                          Sequence 10
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                                             ABR44166 standard;
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9; Conserv
                                                                                                                                                  AVSEIQLMHN
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ABR44166

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Chang M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel fusion peptide comprising self cell-penetrating Tat peptide bound to human parathyroid hormone-derived peptide, useful as component of skin slimming cosmetic composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fusion peptide; tat; hPTHDP; parathyroid hormone; skin; cosmetic;
lipolysis; human; hPTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a fusion peptide (Tat-hPTHDP), where self cell-penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmecic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endothelium does not cause skin disease and has superior lipplysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
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15-MAR-2002; 2002KR-00014062.
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                                                                                                                                                                                                                                    hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                                                                                                                                   Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
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9; Conserv
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Pred. No. 0.061;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel method for determining (M1) the content CC of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CF or diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or typer-parathyroidism. The method (unlike known assays) recognizes that CF of the content of (A) shorter than the complete (64 aa) peptide are active and (ii) that apparently intact peptide may be biologically CF active, and also takes into account the fact that some fragments of (A) care antagonistic (these have the receptor-binding site but lack the N-CF ophysiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      JLT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (IMMU-) IMMUNDIAGNOSTIK AG
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                                                                                                                                                                                                                                               Cyclic peptide; resin bound; parathyroid hormone; osteopathic; disease; hypotensive; PTH receptor; treatment; hyper-calcemia; hypo-calcemia; osteoporosis; osteopenia; hyper-parathyroidism; hypo-parathyroidism; Cushing's syndrome; renal failure; hypertension; bone fracture repair.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 3; 10pp; German.
                                                                                                  Misc-difference
                                                                                                                                  Misc-difference
                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                    Resin bound cyclic peptide 33.
                                                                                                                                                                                                                                                                                                                                                                   09-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY50600 standard; peptide; 11
                                   Misc-difference
                                                                  Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 VSEIQLMHN 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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   /note=
                                                                                                                                                                                   Location/Qualifiers
                                                 /note=
                                                                                  note=
                                                                                                                    note=
                                                                                                                                                   'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.3%;
                   "His(Trt)"
                                                                                                                                                     "FMOC-Ala"
                                                    "Gln(Trt)"
                                                                                                                  "Ser (OtBu) "
                                                                                  "Glu (OtBu) "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 45;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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This invention describes a novel method for the preparation of peptides CC (II) that contain both cyclic and linear peptide fragments comprises CC sequential reaction of a resin-bound linear fragment with the cyclic CC fragment in N-protected form and optionally other linear fragments. The CC products of the invention have osteopathic and hypotensive activity. (II) CC bind to hPTH receptors and act as agonists or antagonists of hPTH. The CC method is particularly used to prepare cyclic peptide analogs of CC parathyroid hormone (PTH) or PTH-related peptides which are useful for CC treating diseases that respond to treatment with agents that bind to PTH CC receptors (with or without activation of adenylyl cyclase activity), e.g. CC hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia of bone fractures. Separate synthesis of the cyclic fragment allows convergent synthesis of resin-bound (II), with better CC preparation of the bridged fragment are confined to a small peptide which can be purified before reaction with the resin-bound component. AAY5568-convergent the peptide fragments described in the method of the
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                                                                   Matches
                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 75; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Convergent synthesis of peptides for treating e.g. bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sledeski AW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-APR-1999;
                                                                                                                                  Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RHON ) RHONE-POULENC RORER PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1999-633822/54.
                                                                   11;
                                                                                   Similarity
                         AVSEIQLMHNLG
AVSEIQL-HNLG
                                                                                                                                      A.
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mencel JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0081897P
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                                                                                   73.7%;
                                                                 Pred. No. 0.2;
0; Mismatches
                                                                                                       DB
                                                                                                       2
                                                                    0
                                                                                                       Length 11;
                                                                      Gaps
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RESULT 10

AAB01862
ID AAB01862;

XX
AC AAB01862;

XX
DT 11-SEP-2000 (first entry)

XX
PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:6.

XX

XX

PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:6.

XX

Calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synth

KW bone synthesis; agonist; osteoporosis; non-parenteral deliv

XX

PM

MO200023594-A1.

XX

PF

27-APR-2000.

XX

20-OCT-1999; 99WO-US024481.
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RESULT 11
AAY96966
ID AAY96
XX
AC AAY96
XX
DT 31-OC
DT 31-OC
XX
KW PTH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         cobiologically active derivatives thereof (AAB01857-B01868). The peptides of the invention are at least 85% identical to the generic peptide of the formula: XI-Val-SerGln-Leu-X3-H18-X4-X5-CC Gly-Lys-X6 (AAB01857) where: XI is Ser or Ala; X2 is 1le or Lys; X3 is CC Met, Leu or N1e; X4 is Asm or Asp; X5 is Leu or Lys; X6 is His or Ser; CC invention also encompass fragments of peptides of the invention of the gravition also encompass fragments of peptides of the invention of some consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- consisting of amino acids 1-9, 1-10, 1-11, 1-12 and cardiovascular systems.

CC and to the recently identified PH-2 receptor. PH has a potent anabolic consisting of process on osteoblasts and renal tubular cells, consisting the receptor againsts of PH-1 and PH-2 receptor. Ahbonlogous calls of process of PH-1 and PH-2 receptor. The peptides of the conditions characterised by a decrease in bone mass, such as conditions characterised by a decrease in bone mass, such as such 
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                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local :
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(POTT/)
(JUEP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Parathyroid hormone (PTH) peptides, PTH related peptides acids that encode them, useful for treating osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel parathyroid hormone (PTH) peptide (AAB01859) and parathyroid hormone-related peptide (PTHIF; AAB01860),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-339693/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-OCT-1998;
PTH; parathyroid hormone; N-terminal; signaling
                                                                             31-OCT-2000
                                                                                                                   AAY96966;
                                                                                                                                                      AAY96966 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GARDELLA T J.
KRONENBERG H M.
POTTS J T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POTTS J T.
JUEPPNER H.
                                                                                                                                                                                                                                                                                                                                    similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ΤJ,
                                                                                                                                                                                                                                                                                                AVSEIQLMH 9
                                                                                                                                                                                                                                                                                                                                      Conservative
                                       hormone N-terminal signaling domain
                                                                             (first entry)
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                                                                                                                                                                                                                                                           9
                                                                                                                                                                                                                                                                                                                            72.9%; Dur
100.0%; Prr
                                                                                                                                                                                                                                                                                                                                                                                                                                                     agonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , MH
                                                                                                                                                                                                                                                                                                                                                       Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Potts JT,
                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                              DB 3; uc.
5. 1.4e+06;
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                                                                                                                                                                                                                                                                                                                                                                            Length 9;
    domain; bone mass,
                                                                                                                                                                                                                                                                                                                                        Indels
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RESULT 12
AAR91645
ID AAR91
XX AAR91
XX AAR91
XX O6-NC
DT O6-NC
CXX Humar
XX Humar
XX Humar
XX Humar
XX Humar
XX OS Synth
XX OE443
XX O4-AH
PN O4-AH
XX O4-SF
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-TP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTH-PTH which avoids the need for regular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
                                                                                                                 Human parathyroid hormone; hPTH; antigen; alpha-helix; diagnosis; active hPTH 1-37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GARD/) GARDELLA T
(KRON/) KRONENBERG
(POTT/) POTTS J T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-DEC-1999;
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                                                                                      Synthetic
                                                                                                                                                             Human parathyroid hormone antigenic peptide hPTH 1-9
                                                                                                                                                                                             06-NOV-1996
                                                                                                                                                                                                                                                       AAR91645 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Page 92; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-452384/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
9; Conserv
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                                                                                                                                                                                                                                                                                                                                              AVSEIQLMH
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                             (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                        72.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                         Score 43;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           1.4
1.4
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                                                                                                                                                                                                                                                                                                                                                                                                         .4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         ø
                                                                                                                                       immunogen;
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28-SEP-1994;

94DE-04434551.

DE4434551-A1

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Best Local S
Matches 8
                                                                                                                           (GARD/)
(KRON/)
(POTT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is a specific example of claimed immunogenic peptides having a sequence from hPTH(1-37) which includes the N-c terminal alpha-helical region and/or the non-structured region for hormone. Antibodies and their binding fragments generated by injenimal with the peptides are useful as diagnostic reagents for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    generated
forms of 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antigenic peptide(s) from generated using them, able to
                                        Parathyroid hormone (PTH) acids that encode them, u
                                                                                            Gardella TJ,
                                                                                                                                                                                                                                                                                                    Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 4; 5pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-180391/19.
                   Disclosure; Page 26;
                                                                                                                                                                       22-OCT-1998;
                                                                                                                                                                                            20-OCT-1999;
                                                                                                                                                                                                                                      WO200023594-A1
                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                     PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:7
                                                                                                                                                                                                                                                                                                                                                           11-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                 AAB01863
                                                                                                                                                                                                                                                                                                                                                                                                   AAB01863 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   determination of biologically active hPTH(\bar{1}-37)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-SEP-1994;
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                                                                        2000-339693/29
                                                                                                                 ) GARDELLA T J.
) KRONENBERG H M.
) POTTS J T.
) JUEPPNER H.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                             Kronenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94DE-04434551
                                                                                                                                                                       98US-0105530P
                                                                                                                                                                                            99WO-US024481.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               67.8%;
88.9%;
                  73pp; English
                                        H) peptides, useful for
                                                                                                                                                                                                                                                                                           osteoporosis;
                                                                                             ×Κ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 40; DB:
Pred. No. 1.4e.
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            distinguish between active and inactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human parathyroid hormone - and antibodies
                                                                                            Potts JT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Maegerlein
                                       PTH related peptides and the nucleic reating osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.4e+06;
0;
                                                                                                                                                                                                                                                                                           TH-2; vitamin D synthesis;
non-parenteral delivery.
                                                                                            Jueppner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         injecting
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The invention relates to

a novel parathyroid hormone

(PTH) peptide

Gardella TJ,

Kronenberg

X X

Potts JT,

Jueppner H;

JUEPPNER H.

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RESULT 14
AAY96981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC biologically active derivatives thereof (AAB01857-B01886) active derivatives thereof (AAB0187-B01886). ARAD1861.

CC generic peptide of the formula: XI Val-Ser-Glu-Ze-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is CC Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; CC provided that the peptide is not PTHTP(1-14). The peptides of the CC invention also encompass fragments of peptides of the invention consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-C terminal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal, cc is the invention of the skeleton, and mediates on osteoblasts and renal tubular cells, CC is the renal and skeleton, and mediates calcium reabsorption, enhances calls of protein brinds to both PTH-1 receptors on osteoblasts and renal tubular cells, CC and is necessary for themson by synthesis in the kidney. A homologous calls of protein brinds to both PTH-1 receptors. PTH has a potent anabolic celfect on the skeleton, and mediates calcium reabsorption, enhances calls of protein brinds to the PTH-2 receptor. PTH has a potent anabolic celfect on the skeletal actions of PTH, and also bind to the PTH-1 creceptor. The peptides of the invention are either agonists of PTH, and also bind to the PTH-1 creceptor. The peptides of the C conditions characterised by a decrease in bone mass, such as content of the pTH-1 receptor. Detectably labelled peptides of the invention are also consisted to a patient of conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via convention are proposed to the corresponding to the corresponding to the corresponding 
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Best Local
                                                                                                                                                                                                                                                                                                                    bone
                                                  (GARD/)
(KRON/)
(POTT/)
                                                                                                                                                                 30-DEC-1999;
                                                                                                                                                                                                     06-JUL-2000
                                                                                                                                                                                                                                       WO200039278-A2
                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                         Parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                 31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY96981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY96981 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9
                                                                                                                             31-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-parenteral route:
PTH-1/PTH-2 receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AAB01859) and parathyroid hormone-related peptide (PTHrP;
                                                                                                                                                                                                                                                                                                                parathyroid hormone; N-terminal; signaling domain; reformation; resorption; remodeling; tetherl; osteo
                                                ) GARDELLA T J.
) KRONENBERG H M.
) POTTS J T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                           hormone N-terminal signaling
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                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                             98US-0114577P
                                                                                                                                                                   99WO-US031108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67.8%;
88.9%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; DB 3; Le.,
; 1.4e+06;
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\begin{array}{c} \mathsf{CCC} \times \mathsf{S} \times \mathsf{S} \times \mathsf{S} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{S} \times \mathsf{S} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{CCC} \\ \mathsf{CCC} \times \mathsf{CCC} \times \mathsf{C
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
                                                                                                                                                                                                                                                                                                  Determining the content of physiologically active parathyroid hormone useful in diagnosis of calcium-metabolism disorders, using two antiboo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                              Disclosure; Page 3; 10pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-376318/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Armbruster FP,
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                                                                                                                                                                                                                                                      reactive with different epitopes.
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88.9%;
                                                                                                                                                         German
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Pred. No. 1.4e+06;
1; Mismatches 0
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This invention describes a novel of active parathyroid hormone (A)

/el method for determining (M1) the content (A) by treating a sample with (i) antibody

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CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CC for diagnosis of calcium-metabolism disorders, osteopathy, or hypocor CC (i) some fragments of (A) shorter than the complete (84 aa) peptide are CC active and (ii) that apparently intact peptide may be biologically CC inactive, and also takes into account the fact that some fragments of (A) care antagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of CC and its 1-37 fragment which may produce falsely high values. This CC sequence represents a peptide fragment used to illustrate the method of CC sequence represents a peptide fragment used to illustrate the method of
                                                          the invention
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В 8 1 AVSEIQLMH SVSEIQLMH Q ω

Matches Query Match

Local Similarity hes 8; Conserv

Conservative

بــر --

Score 40; DB 4; Li Pred. No. 1.4e+06; l; Mismatches 0;

Length 9;

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Gaps

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67.8%;

Sequence 9 AA;

Search completed: May 18, 2004, 10:01:54 Job time : 45.75 secs

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Minimum
Maximum
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Perfect score:
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Match
                                          US-09-730-174A-5
59
1 AVSEIQLMHNLG 1:
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US-09-730-174A-2
US-09-730-174A-3
US-10-168-185-1
US-10-168-185-7
US-10-168-185-7
US-10-192-673-6
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RESULT 2
US-09-730-174A-3
; Sequence 3, Application US/09730174A
; Patent No. US20020110871A1
; GENERAL INFORMATION:
; APPLICANT: Zahradnik, R.J.
; TITLE OF INVENTION: Antibodies and Pei
; TITLE OF INVENTION: Selective Bindin;
; FILE REFERENCE: IMUNE-001A

and Peptide Antigens for Producing Antibodies having Binding Specificity to Bioactive Intact Parathyroid

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4, 10:04:51; Search time 33.25 Seconds (without alignments) 100.425 Million cell updates/sec	24 40.7 10 16 US-10-264-309-369 24 40.7 11 12 US-10-609-217-41 24 40.7 11 12 US-10-632-388-41 24 40.7 11 12 US-10-651-723-41
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/ptodata/1/pubpaa/US07_NEW_PUB.pep:* /ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:* /ptodata/1/pubpaa/US08_NEW_PUB.pep:* /ptodata/1/pubpaa/US08_PUBCOMB.pep:* /ptodata/1/pubpaa/US09A_PUBCOMB.pep:* /ptodata/1/pubpaa/US09A_PUBCOMB.pep:* /ptodata/1/pubpaa/US09B_PUBCOMB.pep:*	
6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:* [6/ptodata/1/pubpaa/US09_EWR PUB.pep:* [6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:* [6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:* [6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:* [6/ptodata/1/pubpaa/US10C_NEW_PUB.pep:* [6/ptodata/1/pubpaa/US10C_NEW_PUB.pep:* [6/ptodata/1/pubpaa/US10C_NEW_PUB.pep:* [6/ptodata/1/pubpaa/US60C_NEW_PUB.pep:*	APPLICANT: Zahradnik, R.J. ; TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having ; TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid; ; FILE REFERENCE: IMUNE-001A ; CURRENT APPLICATION NUMBER: US/09/730,174A ; CURRENT FILING DATE: 2000-12-05 ; NUMBER OF SEQ ID NOS: 12 ; SEQ ID NO 5
er of results predicted by chance to have a requal to the score of the result being printed, the total score distribution.	TYPE: PAT TYPE: PAT ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Induces Formation of Antibodies and Isolates said Antibodies US-09-730-174A-5
DB ID Description	Query Match 100.0%; Score 59; DB 9; Length 12; Best Local Similarity 100.0%; Pred. No. 0.00063; Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
US-09-730-174A-5 Sequence 3 US-09-730-174A-1 Sequence 3 US-09-730-174A-1 Sequence 1 US-09-730-174A-6 Sequence 0 US-09-730-174A-4 Sequence 4 US-09-730-174A-2 Sequence 2 US-10-168-185-7 Sequence 2	Qy
14 US-10-168-185-7 Sequence 7 14 US-10-192-673-6 Sequence 6 14 US-10-158-185-8 Sequence 6 14 US-10-168-185-8 Sequence 7	US-09-730-174A-3 ; Sequence 3, Application US/09730174A ; Patent No. US20020110871A1 : GENERAL TNEORMATION.

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RESULT 3
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                                                                     ; OTHER INFORMATION: Induces Formation of Antibodies and Isolates US-09-730-174A-6
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 3
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 6
                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid is
FILE REFERENCE: IMUNE-001A
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SEQ ID NO 1
LENGTH: 11
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                Query Match
Best Local Similarity
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Best Local
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CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
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TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid;
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                                                                                                                                     LENGTH: 12
TYPE: PRT
                                                                                                               ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Induces formation of Antibodies and Isolates said Antibodies
                                                                                                      FEATURE:
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Score 55; DB 9; Pred. No. 0.0033; 0; Mismatches
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Pred. No.
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US-09-730-174A-2
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US-09-730-174A-4
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US-10-168-185-1
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                                                                                      ; Sequence 1, Application US/10168185; Publication No. US20030175802A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hoselt Reference: IMUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12

LENGTH: 11

TYPE: PRT
ORGANISM: Artificial Sequence
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Patent No. US20020110871A1
GEMERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
APPLICANT: Zahradnik, R.J.
APPLICANT: NVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hoffile REFERENCE: IMUNE-001A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
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SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09730174A Patent No. US20020110871A1
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Best Local
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APPLICANT: Armbruster, Franz Paul APPLICANT: Missbichler, Albert APPLICANT: Schmidt-Gayk, Heinrich APPLICANT: Roth, Heinz-Jurgen TITLE OF INVENTION: Method for Determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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OTHER INFORMATION:
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Parathormone

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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
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US-10-168-185-7
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Best Local Similarity
"Arches 9; Conserve
                                                                                                                                                                                                                                  RESULT 9
US-10-192-673-6
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                                                                                                                                                                 Sequence 6, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
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                    APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Braathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
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REFERENCE: 0609.4570002
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100.0%; Pr
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Pred. No. 0.11
1; Mismatches
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APPLICANT: Anthruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SCOTTWARE: FastSEQ for Windows Version 4.0
                                                           Sequence 7, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Fotts, John T.
APPLICANT: Juppner, Harald
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US-10-168-185-8
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Best Local S
Matches 9
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Publication No. US20030175802A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
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APPLICANT: Juppner, Harald TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
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8; Conserva
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Similarity 100.0%; Pred. No.
9; Conservative 0; Mismatcl
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100.0%; Pred. No.
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1e+06;
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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
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US-10-192-673-8
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                                                                                                               Sequence 8, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
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Best Local
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PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 2000-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
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APPLICANT: Missbichler, Albert
                                     APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
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TYPE: PRT
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8; Conserv
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Pred. No. 1e+06;
1; Mismatches 0
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Pred. No. le+06;
1; Mismatches 0
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Sequence 10, Application No. US20 GENERAL INFORMATION:

Application US/10192673

APPLICANT: Gardella, Thomas J. APPLICANT: Kronenberg, Henry APPLICANT: Potts, John T. APPLICANT: Juppner, Harald

PPLICANT: Juppner, Harald
ITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives

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APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION WIMBER: POT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: US. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
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; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-10-192-673-8
RESULT 15
US-10-192-673-10
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Best Local S
Matches 7
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Publication No. US20030175802A1
                                                                                                                                                                                                                                                           SEQ ID NO 3
LENGTH: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinric
APPLICANT: Roth, Heinz-Jurgen
                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
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7; Conserv
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77.8%;
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Pred. No.
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Pred. No.
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TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609 44570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: US. 09/421,379
PRIOR APPLICATION NUMBER: US. 09/421,379
PRIOR APPLICATION NUMBER: US. 60/105,530
PRIOR APPLICATION NUMBER: US. 60/105,530
PRIOR APPLICATION NUMBER: US. 60/105,530
PRIOR FILING DATE: 1998-10-22
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE: PATURES: PATENTION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: peptide
US-10-192-673-10

Query Match
Best Local Similarity 66.7%; Pred. No. 1e+06;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps

Qy
1 AVSEIQLMH 9
Db 1 SVSEHQLLH 9
Search completed: May 18, 2004, 10:21:00

Search completed: May 18, 2004, 10:21:00
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Run on:

6, Appli
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42, Appli
74, Appl
74, Appl
75, Appl
76, Appl
77, Appl

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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Match
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| Cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
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Patent No. 6563993

GENERAL INFORMATION:
APPLICANT: Sledeski, Adam W.
APPLICANT: Mencel, James J.
TITLE OF INVENTION: PEPTIDES
TITLE OF INVENTION: PEPTIDES
FILE REFERENCE: A31138-US
CURRENT APPLICATION NUMBER: US/09/442,989
CURRENT APPLICATION NUMBER: 60/081,897
EARLIER APPLICATION NUMBER: 60/081,897
EARLIER APPLICATION NUMBER: 61/081,897
EARLIER FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin Ver. 2.1
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (1)
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; NAME/KEY: MOD RES
; LOCATION: (8)
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OTHER INFORMATION:
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US-09-551-976-10
US-09-551-976-10
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US-08-160-604-73
US-08-652-87-71
US-08-160-604-73
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Best Local S
Matches 11
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
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Best Local Similarity 90.0
                                                                 Sequence 6, Application US/09421379
Patent No. 6495662
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/
APPLICATION NUMBER: PCT/EP95/
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 0782
REFERENCE/DOCKET NUMBER: 0782
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: po
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7: USA
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191 Peachtree Street, 37th Floor
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       US-08-817-547A-2
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CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP) FILE REFERENCE: 0609 4570001
                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95,
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 078:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                      MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
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CTTY: Atlanta
CPOTGia
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                    TYPE: amino acid
STRANDEDNESS: unl
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Y: USA
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Pred. No.
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Matches Query Match Best Local

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AVSEHQLEHN 10

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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrp)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
                                          Query Match
Best Local Similarity
Marches 8; Conserva
                                                                                                                                                                                                                                                                                                                HYPERCALCEMIA OF MALIGNANCY-PTHRP
HUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/715,280
                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Patent No. 5460978; APPLICANT: MARTIN, THOMAS J.; MOSELEY, JANE M.; KEMP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-421-379-7
                                                                                                                                                                                                                                                                                                                                                                                                  BRUCE E.; WETTENHALL, RICHARD E.H.
TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 8
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SEQ ID NO 7
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7, Application US/09421379 Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/
FILING DATE: 14-JUN-1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                      APPLICATION NUMBER: 199,235 FILING DATE: 09-MAY-1988 APPLICATION NUMBER:
                                                                                                                                                               LENGTH: 11
                                                                                                                                                                                                   FILING DATE:
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    AVSEIQLMHN 10
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80.0%;
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88.9%;
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                                                            Score 35; DB of Pred. No. 2.2; O; Mismatches
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Pred. No. 3e+05;
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                                                                               DB 6;
2.2;
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APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: October, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Hormone-related Peptide (PTHP) and Parathyroid
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION UNUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
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Patent No. 6030790
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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ADDRESSEE: Jones & P
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Peptides from the hPTH Sequence NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                          CITY: Atlanta
STATE: Georgia
                                                                                                                                                               FILING DATE:
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7; Conserv
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191 Peachtree Street,
                                                                                                                                                                                                                                                                                                                                          USA
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77.8%;
                                                                                                                                                                                                                      Release #1.0, Version
                                                                                                                       PCT/EP95/03757
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                       07826-0007
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Pred. No. 3e+05;
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TYPE: PRT

5460978-3

SEQ ID NO:3:

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APPLICANT: Kronenberg, Henry
APPLICANT: potts, John T.
APPLICANT: Duppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.0
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US-08-817-547A-4
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OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-09-421-379-10
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                                                                                                                                         Sequence 4, Application Patent No. 6030790 GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides f:
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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STRANDEDNESS: un
ADDRESSEE: Jones & Askew, LLP STREET: 191 Peachtree Street, 37th Floor
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                                                                                                                                                                                                                                                                                                                                        50.8%;
Similarity 66.7%;
6; Conservation
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87.5%;
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Pred. No. 3e+0
1; Mismatches
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Pred. No. 3e+05;
                                                                        from
                                                                      the hPTH Sequence
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US-08-817-547A-4
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Patent No. 5885808
APPLICATION NUMBER: US/
FILING DATE: 07/05/95
CLASSIFICATION: 514
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Jules E. Goldberg
STREET: 261 Madison Avenue
                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                             APPLICANT: Spooner, Robert A.
APPLICANT: Epenetos, A.A.
TITLE OF INVENTION: Compounds to target cells
NUMBER OF SEQUENCES: 80
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MOLECULE TYPE:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STRANDEDNESS: uni
                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 10016-2391
                                                                                                                                                                                                                                                                             CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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85.7%;
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22,176
07826-0007
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                                                                                               US/08/428,257A
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                                            14:
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                                                                                                                                   (EPO)
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; TOPOLOGY: 1: ; MOLECULE TYPE: US-08-428-257A-14

protein

TYPE:

amino acid OGY: linear

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RESULT 12
US-08-802-981-124
                                                                                                             RESULT 13
US-08-177-109A-56
                                                                                                                                                                                                                                                                                                                              ) OTHER INFORMATION: /product= "Aib" US-08-802-981-124
Sequence 56, Application US/08177109A
Patent No. 5869615
GENERAL INFORMATION:
APPLICANT: Dennis E. Hourcade and
TITLE OF INVENTION: MODIFIED COMP.
NUMBER OF SEQUENCES: 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/802,981
FILING DATE: 20-FEB-1997
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6037137
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGE... 10m NAME: Hutter, Tom NAME: Hutter, Tom NAMER: 38,498 REGISTRATION NUMBER: 0168 REFERENCE/DOCKET NUMBER: 0168 TELECOMMUNICATION INFORMATION: 125,576,0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 576-030 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NEARL INFORMATION: Komoriya, Akira
APPLICANT: Komoriya, Akira
APPLICANT: Packard, Beverly S.
TITLE OF INVENTION: Compositions for the Detection of Enzyme
TITLE OF INVENTION: Activity in Biological Samples and Methods of Use Thereof
TITLE OF SECUENCES: 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Two Embarcac
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH:
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6037137
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83.3%;
                     . Hourcade and Teresa J. Oglesby MODIFIED COMPLEMENT PROTEASES
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Pred. No. 1.4e+02;
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Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                           Mismatches
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HYPOTHETICAL:
US-08-177-109A-56
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Best Local Similarity
3; Conserva
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NUMBER OF SEQUENCES: 62

CORRESPONDENCE ADDRESS: Patrea L. Pabst STREET: 2800 One Atlantic Center STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                    US-08-687-706-56
Patent No. 5928892
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/177,109A
FILING DATE: 03-JAN-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION UNMBER: 31.284
REFERENCE/DOCKET NUMBER: WU 107
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 56,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (404) F
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ZIP: 30309-3450
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Y: USA
                                                                                                                                                                                                                                                                                                                                                                                     Application US/08687706
                                                                                                                                                                                                                                                                                                                                 Dennis E. Hourcade and Teresa VENTION: MODIFIED COMPLEMENT F
                                                                                                                                                                                                                   USA
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Pred. No. 3e+05;
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                                                                                                                                                                                                                                                                                                                                  PROTEASES
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COMPUTER READABLE FORM:

MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
APPLICATION NUMBER: US/08/817,547A
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 07826-0007
TELEPAXI: 404-818-3700
TELEPAX: 404-818-3700
TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acids
TYPE: amino acids
STRANDEDNESS: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: no
US-08-817-547A-17
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Patent No. 6030790
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Hock, Dieter
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 40.7%;
Best Local Similarity 60.0%;
Matches 3; Conservative
Query Match 40.7%; Score 24; DB 3; Best Local Similarity 100.0%; Pred. No. 3e+05; Matches 4; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPAX: (404) 873-8795
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
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REFERENCE/DOCKET NUMBER: WU 107 DIV
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 MHNLG 12
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1 LHNMG 5
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Pred. No. 3e+05;
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                                     Length 5;
     0; Indels
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Qy 9 HNLG 12
Db 1 HNLG 4
Search completed: May 18, 20

Search completed: May 18, 2004, 10:06:44 Job time : 12.75 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score
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Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               May 18, 2004, 09:56:10 ; Search time 10.25 Seconds (without alignments)
112.514 Million cell updates/sec
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61
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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29.5
27.9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length DB
                                                                       GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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E49033
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S33844
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S21205
                                                                       A38841
A55837
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S51737
A35779
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                                                                                                                                                                                                                                    amine oxidase (cop
polyphosphate-gluc
Ig heavy chain CDR
N-methylpurine DNA
alpha-2-macroglobu
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                                                                                       T-cell receptor al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carnocin UI49 - Carnocin UI49 
                    ribosomal protein hypertrehalosemic
                                                                  neuropeptide Antho
                                                                                                                                          5-aminoimidazole r
                                                                                                                                                                                     T-cell receptor ga
                                                                                                                                                                                                                                                                                                                                                               hypertrehalosemic
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hypertrehalosemic
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Best Local S
Matches
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Best Local :
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cytochrome b559 component psbF - pepper chloroplast (fragment)
C;Species: chloroplast Capsicum annuum (pepper)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 11-Jun-1999
C;Accession: S28055
R;Kuntz, M.; Camara, B.; Weil, J.H.; Schantz, R.
Plant Mol. Biol. 20, 1185-1188, 1992
A;Title: The psbt gene from bell pepper (Capsicum annuum): plastid RNA editi
A;Reference number: S28055; MUID: 93099270; PMID: 1463853
A;Accession: S28055; MUID: 93099270; PMID: 1463853
                                                                                                                                                                                                                                                                       RESULT 2
PHILE7
C. Species: Homo sapiens (man)
C. Species: J. L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Wideling (man)
C. Species: J. Species: Homo sapiens (man)
A; Tile: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor
A; Reference number: S26512; MUID:92364546; PMID:1380061
A; Accession: PHILE7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Genome: chloroplast
C;Superfamily: cytochrome b559 component
C;Keywords: chloroplast; photosynthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Molecule type: DNA
A;Residues: 1-10 <KUN>
A;Ccsos-references: EMBL:X65570; NID:g14344; PIDN:CAA46539.1; PID:g415734
C;Genetics:
                                                                                                                                                                                                 A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
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3; Conserve
                                                                                               Similarity
5; Conserv
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AVSETGE
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37.5%; Pred. No. 7.4e+0:
tive 4; Mismatches
                                                                                                                          32.8%;
                                                                                                 Score 20; DB 2;
Pred. No. 9.1e+02;
0; Mismatches 2
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photosystem II; thylakoid
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A; Residues:
C; Keyword
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A;Title: Purification and characterization of a new bacteriocin isolated from a Carnobad A;Reference number: A58718; MUID:92321768; PMID:1622206
A;Accession: A58718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998
C;Accession: S78420
R;Goldschmidt-Reisin, S.; Graack, H.R.
Submitted to the Protein Sequence Database, Februar
Вb
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A; Accession: $78420
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C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change
C;Accession: A58718
                               δÃ
                                                                                                                                                                                                                                                                      C;Accession: A60589
R;Yoshino, K.I.; Kajiura, H.;
Comp. Biochem. Physiol. B 94,
                                                                                                                                                                                                                                                                                                                     sperm-activating peptide (Tyr-2, Asn-3, Gly-5, Ile-9, Asp-10 SAP-I) - slate-pencil urchi
C,Species: Heterocentrotus mamillatus
C,Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
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                                                                                                                                                               A;Molecule type: protein A;Residues: 1-10 < YOS>
                                                                                                                                                                                              A; Accession: A60589
                                                                                                                                                                                                                      A, Reference number: A60527
                                                                                                                                                                                                                                   A;Title: A halogenated amino acid-containing sperm activating peptide and its related peotus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
                                                                                                                                                                                                                                                                                                                                                                                      A60589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ribosomal protein RL41, mitochondrial [validated] - rat (fragment)
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Matches
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                                                                                                                                            ;Superfamily: unassigned animal
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Best Local :
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 YNLG 5
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75.0%;
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                                                                      Score 18; DB 2; Pred. No. 1.8e+03; Mismatches 0
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Pred. No. 2.8e+05;
0; Mismatches 1
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R,Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R.
J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences and antibody properties of anti-
A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: A37268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rinehart Jr., Biochem. Biophys. Res. Commun. 124, 350-358, 1984
A;Title: Structures of two cockroach neuropeptides assigned by fast atom bombardment A;Reference number: A90118; MUID:85046530; PMID:6548628
A;Accession: A05169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neuropeptide M-I - American cockroach (Species: Periplaneta americana (American cockroach) (Species: Periplaneta americana (American cockroach) (Spate: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 31-Dec-1993 (Accession: A0516) (A
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A05169
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C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 8
A37268
A37268
Ig heavy chain C region (129) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 13-Mar-1998 #text_change
C;Accession: A37268
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A; Residues: 1-12
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J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for
A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1190
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Residues: 1-8 <WIT>
C; Keywords: neuropeptide
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                                                                                                                                                                                                                                 A;Status: preliminary
A;Molecule type: mRNA
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                                                                Score 17; DB 2;
Pred. No. 2.9e+03;
1; Mismatches 4
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Pred. No. 2.8e+05;
1; Mismatches 3
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ESQSFPNVG

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A; Molecule type: protein
A; Residues: 1-12 < SCH>
A; Residues: 1-20 < SCH>
A; Residues: 1-10 < SCH>
A; Note: the amino end of this peptide is not blocked
A; Note: synthetic locustamyotropin mimics natural locustamyotropin only in the C; Comment: This peptide was shown to stimulate visceral muscle contractions in C; Superfamily: pyrokinin
C; Superfamily: pyrokinin
C; Superfamily: pyrokinin
C; Keywords: amidated carboxyl end; neuropeptide
F; 12/Modified site: amidated carboxyl end (Leu) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Makiya, R.; Stigbrand, T.
Eur. J. Biochem. 205, 341-345, 1992
A;Title: Placental alkaline phosphat
A;Reference number: S21205; MUID:922
A;Accession: S21205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 16-Aug-1996
C;Accession: S21205
                                                              A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
                                                                                                                         C;Accession: PH1189

R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, P. J. Exp. Med. 176, 439-447, 1992

A;Title: H-2-restricted cytolytic T lymphocytes specific A;Reference number: S26512; MUID:92364546; PMID:1380061

A;Accession: PH1189
                                                                                                                                                                                                                                                     T-cell receptor alpha chain V region (Cw3/2C3) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-12 <MAK>
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A;Title: Isolation, identification and synthesis of locustamyotropin A;Reference number: A43975; MUID:90341077; PMID:1974346
A;Accession: A43975
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C;Species: Locusta migratoria (migratory locust)
C;Date: 11-Feb-1993 #sequence_revision 02-Jun-1994 #text_change 08-Dec-1995
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Pred. No. 3.5e
3; Mismatches
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C;Species: Carausius morosus
C;Date: 30-Sep-1993 #sequence revision 30-Sep-1993 #text_change
C;Accession. UC1416; S07157
R;Gaede, G.; Kellner, R.; Rinehart, K.L.; Proefke, M.L.
Biophys. Res. Commun. 189, 1303-1309, 1992
A;Title: A tryptophan-substituted member of the AKH/RPCH family
A;Reference number: JC1416; MUID:93129188; PMID:1482345
A;Accession: JC1416
                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Residues: 1-10 <GABZ1>
R;Residues: 1-10 <GABZ1>
R;Gaede, G.; Rinehart Jr., K.L.
Biol. Chem. Hoppe-Seyler 368, 67-75, 1987
Biol. Chem. Hoppe-Seyler of the hypertrehalosaemic factor II from the corpus A;Title: Primary structure of the hypertrehalosaemic factor II from the corpus A;Reference number: S07157; MJID:87157103; PMID:3828078
A;Accession: S07157
A;Molecule type: protein
A;Residues: 'Z',2-10 <GABZ2
A;Residues: 'Z',2-10 <GABZ2
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                                                                                                                                                  C;Keywords: amidated carboxyl end; glycoprotein; hormone; neuropeptide; p F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;8/Binding site: carbohydrate (Trp) (covalent) #status experimental F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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C;Species: Cryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 05-Nov-1999
C;Accession: 146868
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                                                                                                                                                                                                                                                C;Comment: Hypertrehalosemic factor II lacks the tryptophan modification.
C;Comment: This peptide raises hemolymph levels of trehalose in the cockroach Periplanets
C;Superfamily: adipokinetic hormone
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A; Residues: 1-7 <FRI>
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Pred. No. 2.8e+05;
1; Mismatches. 1
                                                      Score 16; DB
Pred. No. 4.5e
2; Mismatches
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amine oxidase (copper-containing) (EC 1.4.3.6) - pig (fragment) C;Species: Sus scrofa domestica (domestic pig) C;Date: 12-Oct-1989 #sequence_revision 31-Dec-1993 #text_change 06-Sep-1996 C;Accession: A32428 R;van der Meer, R.A.; van Wassenaar, P.D.; van Brouwershaven, J.H.; Duine, J.A. Biophys. Res. Commun. 159, 726-733, 1989 A;Title: Primary structure of a pyrroloquinoline quinone (PQQ) containing pepti A;Reference number: A32428; MUID:89193662; PMID:2539124 A;Accession: A32428
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A32428
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N;Alternate names: Cam.-HrTH-II
C;Species: Extatosoma tiaratum
C;Date: 30-Jun-1992 #sequence_revision 24-Oct-1997 #text_change 31-Oct-1997
C;Accession: S09138
R;Gaede, G; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
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A;Accession: 809138
Search completed: May 18, 2004, 10:05:41 Job time: 10.25 secs
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A; Residues: 1-7, 'K', 9-11 < VAN>
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ALIGNMENTS	FIBB_ERYPA	HTF2_PERAM	ANG2_BOTJA	RE31_LITRU	RE21_LITRU	UKA2_HUMAN	Q2OA_COMTE	PVK1_PERAM	CSI5_BACSU	CORZ PERAM	ASL2_BACSE	PPK1_PERAM
			Q10582 bothrops ja	P82072 litoria rub	P82071 litoria rub	P31144 homo sapien	P80464 comamonas t	P41837 periplaneta	P81095 bacillus su	P11496 periplaneta	P83147 bacteroides	P82691 periplaneta

RESULT 1 PSBF_CAPAN PSBF_CAPAN Q03367; 01-JUN-1994 EMBL; X65570; CAA46539.1; ... PIR; S28055, S28055. HAMAP; MF_00643; -; 1. InterPro; IPR006216; Cyt_b559. PROSITE; PS00537; CYTOCHROME_B559; PARTIAL. Chloroplast; Photosystem II; Heme; Electron transport; Transmembrane. NON_TER. 1 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch). Kuntz M., Camara B., Weil J.-H., Schantz R.; "The psbL gene from bell pepper (Capsicum annum): plastid RNA editing also occurs in non-photosynthetic chromoplasts."; plant Mol. Biol. 20:1185-1188(1992). -!- FUNCTION: This b-type cytochrome is tightly associated with th reaction center of photosystem II and possibly is part of the water-oxidation complex. -!- SUBUNIT: Heterodimer of an alpha subunit and a beta subunit. -!- SIMILARITY: Belongs to the psbE / psbF family. Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamiids; Solanales; Solanaceae; Capsicum. STRAIN=cv. Lamuyo; TISSUE=Fruit, MEDLINE=93099270; PubMed=1463853; Chloropiast. 01-JUN-1994 10-OCT-2003 SEQUENCE FROM N.A. NCBI_TaxID=4072; Capsicum annuum (Bell pepper). (Fragment). 4 (Rel. 29, Created) 4 (Rel. 29, Last sequence update) 3 (Rel. 42, Last amoutation update) 5559 beta subunit (PSII reaction center subunit VI) STANDARD; and Leaf; 10 B the

1 SISAMQFI 1 AVSEIQEM

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32.8%; milarity 37.5%; Conservative

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01-JUN-1994
01-JUN-1994
16-OCT-2001
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MOD_RES 1 1 PYRR
MOD_RES 10 10 AN; 1163 MW; 0567
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Appl. Environ. Microbiol. 58:1417-142
-!- FUNCTION: Lanthionine-containing:
Active on Gram-positive bacteria.
Antibictic; Bacteriocin; Lantibiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-89145002; PubMed=3226948;
Gaede G., Hilbich C., Beyreuther K., Rinehart K.L. Jr.;
"Sequence analyses of two neuropeptides of the AKH/RPCH-family
the lubber grasshopper, Romalea microptera.";
Peptides 9:681-688(1988).
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that
                                                                                                                                                                                                                                                              -!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose is major carbohydrate in the hemolymph of insects).
|- SUBCELLULAR LOCATION: Secreted.
|- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
|- InterPro; IPR002047; AKH.
|- InterPro; IPR002047; AKH.
|- PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Romalea microptera (Lubber grasshopper).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Romaleidae; Romalea.
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TISSUE=Corpora cardiaca;
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Ol-JUN-1994 (Rel. 29, Last sequence uj
16-OCT-2001 (Rel. 40, Last annotation
Lantibiotic carnocin UI49 (Fragment).
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(Hypertrehalosaemic factor).
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en-Meyer J., Gudmundsdottir
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30-MAY-2000
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                                                                                            Duve H., Johnsen A.H., Maestro J.-L., Scott Davey M., East P.D., Thorpe A.; "Lepidopteran peptides of the allatostatin @ Peptides 18:1301-1309(1997).

-!- SIMILARITY: Belongs to the allatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide; Amidation; Pyrokinin
MOD_RES 12 12 AMIDA
SEQUENCE 12 AA; 1213 MW; D766C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schoofs L., Holman G.M., Hayes T.K., Tips A., Nachman R.J., Vandesande F., de Loof A.; Vandesande F., de Loof A.; "Isolation, identification and synthesis of locustamyotropin (Lom-MT), a novel biologically active insect peptide."; Peptides 11:427-433(1990).

-!- FUNCTION: Mediates visceral muscle contractile activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Locusta migratoria (Migratory locust).
Eukaryota; Metazoa; Arthropoda; Haxapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-AUG-1991 (Rel. 19, Created
01-AUG-1991 (Rel. 19, Last so
01-AUG-1991 (Rel. 19, Last so
01-AUG-1991 (Rel. 19, Last so
Locustamyotropin 1 (LOM-MT-1)
                                 Neuropeptide; Amidation. MOD_RES 8 8
                                                                                                                                                                                                                                     MEDLINE=98054539;
Duve H., Johnsen F
                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                             Neoptera; Endopterygota; Lepidoptera;
Tortricoidea; Tortricidae; Olethreuti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cydiastatin
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P82157;
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                                                                                                                                                                                                                                                                                                               TISSUE=Larva;
                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=82600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cydia pomonella (Codling moth)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR001484; Pyrokinin.
PROSITE; PS00539; PYROKININ; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE.
TISSUE=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=7004;
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PIR; A43975; A43975.
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Schoofs L., Holman G.M., Hayes T.K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              activity)
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(Rel. 39,
(Rel. 39,
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                                                                                                                                                                                                                                 PubMed=9392829;
1.H., Maestro J.-L.,
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annotation
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Pred. No. 2.
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D766C92722D6DDDD
                                     AMIDATION
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SEQUENCE

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936 MW;

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RESULT 6
HTF2 CARMO
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Best Local
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                                                                                                                                                                                                                                                                                                                                         Neuropeptide; Amidation; Glycoprotein; Pyrrolidone carboxylic MOD RES
1 1 PYRROLLDONE CARBOXYLIC ACID.
CARBOHYD 8 8 C-LINKED (MAN) (PROBABLE).
MOD RES
10 10 AMIDATION.
SEQUENCE 10 AA; 1164 MW; 9B9036745771A9D1 CRC64;
                                                                                                                                                                                                                                                                                                       PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the corpora cardiaca of the cockroaches Leucophaea maderae, Gromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry."; Blol. Chem. Hoppe-Seyler 371:345-354(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaede G., Rinehart K.L. Jr.;
"Primary structures of hypertrehalosaemic neuropeptides isolated from the corpora cardiaca of the cockroaches Leucophaea maderae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Primary structure of the hypertrehalosaemic factor II from the corpus cardiacum of the Indian stick insect, Carausius morosus, determined by fast atom bombardment mass spectrometry."; Biol. Chem. Hoppe-Seyler 368:67-75(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=C.morosus; TISSUE=Corpora cardiaca; MEDLINE=87157103; PubMed=3828078; Gaede G., Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Extatosoma tiaratum (Stick insect).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Phasmatodea; Euphasmida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUI-1989 (Rel. 11, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-CCT-2003 (Rel. 42, Last annotation updat
Hypertrehalosaemic factor II (HTF-II) (HRTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPECIES=E.tiaratum; TISSUE=Corpora MEDLINE=90253659; PubMed=2340112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neuropeptide II)
Carausius morosus (Indian stick insect),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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Heteronemiidae; Carausius
                                                                                                                                                                                                                                                          PROSITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CARBOHYDRATE-LINKAGE SITE.
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                                                                                                                                                                                                                                                                                                  JC1416; JC1416.
S09138; S09138.
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  EIQFMHNLG
                                                                                                                                                                                                                                                        PS00256;
                                                                                                                                                                                                                                                                              IPR002047; AKH.
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                                                                                                                                                                                                                                                        AKH; 1.
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Pred. No.
                                              Score 16; DB Pred. No. 2.9e 2; Mismatches
                                                 2;
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Phasmatoidea;
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E103 L1
AC P82059
DT 28-FEB
DT 10-OCT
DE Electr
OS Litori
OC Amphib
OC Pelodr
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CX NCBI 
                                                                                                                                                                                                                                                                              SO CEL PROCESS
RESULT 9
EFG_CLOPA
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Best Local :
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P81010;
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Amphibia; Batrachia; An
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P82099;
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Submitted (JUL-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUSSO
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10-OCT-2003
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                                                                                                                                                                                                                                                                                SEQUENCE
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2; Conserv
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15-UUL-1998 (Rel. 36, Created)
15-UUL-1998 (Rel. 36, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
Allergen Fus s 13596* (Fragment).
Fusarium solani (subsp. pisi) (Nectria haem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Skin secretion;
Wabnitz P.A., Bowie J.H., Tyler M.J.,
West from the skin glands of the
Litori electrica. Comparison with the
                                                                                                                                                                                               Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordari
Hypocreomycetidae; Hypocreales; Nectriaceae; Nectria.
NCBI_TaxID=70791;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amphibian defense peptide; Amidation MOD_RES 5 AMIDATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Litoria rubella (Desert tree frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION: Sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Rel. 41, Created)
(Rel. 41, Last sequence update)
(Rel. 42, Last annotation update)
                                                                                                                                        TISSUE=Mycelium;
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Anura; Neobatrachia; Hyloidea;
             24.6%;
                                                                                                      Swiss-Prot.
allergic re
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                                                            C372C441F5B69041 CRC64;
               Score 15;
Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                     (Nectria haematococca)
 Mismatches
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                                                                                                          reaction
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Australian buzzing tree frog
skin peptides from Litoria
                                                                                                                                                                                                                                                                                    update)
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1.4e+05;
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Hylidae;
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RESULT 10
NUHM_CANFA
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Best Local S
Matches 2
                               MEDLINE-98163340; PubMed-9504812;

A Dunn M.J., Corbett J.M., Wheeler C.H.;

PUNCTION: TRANSFER OF ELECTRON SEROM NADH TO THE RESPIRATORY

CHAIN. THE INMEDIATE ELECTRON ACCEPTOR FOR THE ENZYME IS BELIEVED

CO BE UBIQUINONE. COMPONENT OF THE FLAVOPROTEIN-SULFUR (FP)

CO BE UBIQUINONE. COMPONENT OF THE FLAVOPROTEIN-SULFUR (FP)

CO BE UBIQUINONE. THE MADH + ubiquinone = NAD(+) + ubiquinol.

COPACTOR: Binds 1 2Fe-28 Cluster (Potential).

-!- SUBUNIT: NADH + acceptor = NAD(+) + reduced acceptor.

-!- SUBUNIT: NADH + acceptor = NAD(+) + SUBUNIT: NADH + SUBSECTIONS | SUBSECTIONS
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P81350;
P81350;
15-JUL-1998
15-JUL-1998
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the GTP-bindir
EF-G/EF-2 subfamily.
InterPro; IPR000795; EF GTPbind.
PROSITE; PS00301; EFACTOR GTP; PARTIAL.
Elongation factor; Protein biosynthesis;
NON_TER 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Flengsrud R., Skjeldal L.;
"Two-dimensional gel electrophoresis separation and N-sequence analysis of proteins from Clostridium pasteur Electrophoresis 19:802-806(1998).

-!- FUNCTION: This protein promotes the GTP-dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUHM CANFA
P49820;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-1996 (Rel. 34, Created)
15-UTL-1998 (Rel. 36, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
NADH-ubiquinone oxidoreductase 24 kDa subunit
(EC 1.6.99.3) (Fragment).
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                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                      Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
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  SIMILARITY:
                    mitochondrial
                                                                                                                                                                                                                                                                                                                                                                     _TaxID=9615;
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Belongs to the
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. 41, Last
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                    membrane.
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Pred. No. 5.1e
3; Mismatches
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complex I
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pasteurianum W5
  subunit
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Canis.
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RESULT
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Matches 2
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Best Local &
Matches
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30-MAY-2000
30-MAY-2000
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MOD_RES
SEQUENCE
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MEDILINE=90319122; PubMed=1973541;

Grimmelikhuijzen C.J.P., Rinehart K.L. Jr., Jacob E., Graff D.,

Reinscheid R.K., Nothacker H.-P., Staley A.L.;

"Isolation of L-3-phenyllactyl-Leu-Arg-Asn-NH2 (Antho-RNamide),
anemone neuropeptide containing an unusual amino-terminal block:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anthopleura elegantissima (Sea anemone)
Eukaryota; Metazoa; Cnidaria; Anthozoa;
Nynantheae; Actiniidae; Anthopleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FLRN AN
P58707;
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NON_TER 11
SEQUENCE 11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    group.";
Proc. Natl.
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10-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                  ALL7_CYDPO
P82158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=6110;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-2003
                                                                                                                                               Eukaryota; Metazoa; Arthropoda; Hexapoda; Insec
Neoptera; Endopterygota; Lepidoptera; Glossata;
Tortricoidea; Tortricidae; Olethreutinae; Cydia
                                                              SEQUENCE
                                                                                                                                                                                                                                   Cydia pomonella (Codling
                                                                                                                                                                                                                                                                  Cydiastatin
                                                                                                                   NCBI_TaxID=82600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C. Natl. Acad. Sci. U.S
SUBCELLULAR LOCATION:
TISSUB SPECIFICITY: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MASS SPECTROMETRY: MW=549.3;
A35779; A35779.
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1 (Rel. 41,
3 (Rel. 42,
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(Rel.
(Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;

    Cmplx1 24kDa.
COMPLEXI 24K; PARTIAL.
    Dbiquinone; Mitochondrion; Metal-binding;
    2Fe-2S.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cnidaria; Anthozoa;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neuron specific.
MW=549.3; METHOD=FAB
                                                                                                                                                                                                                                      moth)
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AMIDATION.
64540729A0000000 CRC64;
                                                                                                                                                                                                                                                                                                 annotation
                                                                                                                                                                                                                                                                                                                             sequence
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1; Mismatches
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Pred. No. 1.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         267F5369C9C72DD8
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                                                                                                                                                                                                          Hexapoda; Insecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                              update)
on update)
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TISSUE=Larva; MEDLINE=98054539;

PubMed=9392829

SORRIBRA

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RESULT 13
TYSI_LITRU STANDARD; PRT; 7 AA.
AC P82065;
DT 28.FEB-2003 (Rel. 41, Last sequence update)
DT 10.CCT-2003 (Rel. 42, Last annotation update)
DT 10.CCT-2003 (Rel. 42, Last annotation update)
DT 10.CCT-2003 (Rel. 42, Last annotation update)
DT 11.CCT-2003 (Rel. 42, Last annotation update)
DT 12.CCT-2003 (Rel. 42, Last annotation update)
DT 12.CCT-2003 (Rel. 42, Last annotation update)
DT 13.CCT-2003 (Rel. 42, Last annotation update)
DT 14.CCT-2003 (Rel. 42, Last annotation update)
DT 15.CCT-2003 (Rel. 42, Last annotation update)
DE 17.CCT-2003 (Rel. 42, Last annotation update)
DE 17.CCT-2003 (Rel. 42, Last annotation update)
DE 17.CCT-2003 (Rel. 42, Last annotation update)
CC NCBLTANDALD MASS SPECTROMETRY.
R PSECUENCE, AND MASS SPECTROMETRY.
R PSECUENCE, AND MASS SPECTROMETRY.
R PSECUENCE FOR UPDATE SPECIFIC P.A., Waugh R.J., Both SECTION UPDATED.
R PSECUENCE AND MASS SPECTROMETRY.
R PSECUENCE AND R PSECUENCE CARBO.
R PYTROLIDONE CARBO.
R MOD RES AND MASS SPECTROMETRY.
R MASS SPECTROMETRY
ALIS CARMA
ID ALIS C
AC P8187
DT 30-MAY
DT 30-MAY
DT CARCIN
OS CAR
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Best Local S
Matches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                               AL18 CARMA
P81821;
30-MAY-2000
30-MAY-2000
30-MAY-2000
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Litoria rubella (Desert tree frog).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;

Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;

pelodryadinae; Litoria.

MCBI TaxID=104895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstan. Davey M., East P.D., Thorpe A., "Lepidopteran peptides of the allatostatin superfamily."; Peptides 18:1301-1309(1997).
-!- SIMILARITY: Belongs to the allatostatin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide;
                                 Eukaryota; Metazoa; Arthropoda;
Eumalacostraca; Eucarida; Decapo
Eubrachyura; Portunoidea; Portun
NCBI_TaxID=6759;
                                                                                                              Carcinus maenas (Common shore crab) (Green crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu
                                                                                                                                                                                                                 Carcinustatin 18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amphibian defense peptide; Amidation; Neuropeptide; Pyrrolidone carboxylic acid.
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2; Conserv
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2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QIPWFH 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EIQFMH 9
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                                                                                                                                                                                                                                               (Rel. 39, Created)
(Rel. 39, Last sequence up
(Rel. 39, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amidation.
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                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
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                                                                                                                                                                                                                                                                                           sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14; DB Pred. No. 1.4e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14; DB 1;
Pred. No. 1.4e+05;
2; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AMIDATION
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TISSUE-Cerebral ganglion, and Thoracic ganglion,
MEDLINE=98121193; PubMed=9461295;
Therean A H. Maestro J.-L., Scott A.G.
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Eur. J. Blochem. 250:727-734(1997).
Eur. FUNCTION: May act as a neurotransmitter
i- SIMILARITY. Belongs to the allatostatin
                                                                                                                                                                                                                                                                                                             MEDLINE=98054539; PubMed=9392829;
Duve H., Johnsen A.H., Maestro J.-L., Scott
Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin s
Peptides 18:1301-1309(1997).
-!- SIMILARITY: Belongs to the allatostatin
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;

Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;

Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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"Isolation and identification of multiple neuropeptides of
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Fibrinolytic metalloproteinase (Fragment).
Agkistrodon contortrix.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=91378546; PubMed=1898066;
Guan A.L., Retrios A.D., Henderson G.N., Markland F.S.Jr.;
"Purification and characterization of a fibrinolytic enzyme
of the southern copperhead snake (Agkistrodon contortrix
contortrix).";
Arch. Biochem. Biophys. 289:197-207(1991).
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NCBI_TaxID=8720;
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Brouillette J.A.,
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Mammalia; Eutheria; Carnivora;
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Canine distemper virus.
Viruses; ssRNA negative-strand
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Liermann H., Harder T., Haas
"Genetic analysis of the cen
proximal coding part of the
morbilliviruses.";
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A
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73; AAF20918.1; -.
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ba; Chordata;
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he central
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STRALN=Dog #10757/96;
Liermann H., Harder T., Haas L.;
Liermann H., Harder T. Haas L.;
"Genetic analysis of the central untranslated genome region "genetic analysis of the F gene of wild-type and vaccine morbilliviruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
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01-JAN-1998 (TrEMBLrel. 05,
01-NOV-1999 (TrEMBLrel. 12,
01-OCT-2002 (TrEMBLrel. 22,
ORF YNL337W (Fragment).
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Raftery M.J., Bradford A.M., Bowie J.H., '
"Peptides from Australian frogs. The stru-
the banjo frogs Limnodynastes interioras,
Limnodynastes terraereginge.";
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachi
Limnodynastinae; Limnodynastes.
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                                                                                                                                                                             Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes.
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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12 AA; 1236 MW; 147AA70FDF472724 CRC64;
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EMBL; AF384567; AAM50405.1; -. GO; GO:0009507; C:chloroplast; I
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EMBL; KO
   Sutrave P., Jansen H.W., Bister K., Rapi
"3'-terminal region of avian carcinoma vi
elements with avian sarcoma viruses Y73
. Virol. 52:703-705(1984).
EMBL; K03100; hAA42388.1; -.
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                                                                                                                                             TISSUE-SKIN SECRETION;
Bradford A.M., Raftery M.J., Bowie J.H., Wa
"Peptides from Australian frogs. The struct
Limnodynastes salmini and Fletcherin from L
Aust. J. Chem. 46:L235-L244(1993).
-i-MASS SECTROMETRY: WM=1114, METHOD=FAB.
SEQUENCE 12 AA; 1114 MW; 3AB5A976CAA727
                                                                                                                                                                                                                                                                                                                                                                                                                                 P82085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Actin multigene family of Dictyostelium. Cold Spring Harb. Symp. Quant. Biol. 46:4 EMBL; K02957; AAA33150.1; JOINED. EMBL; K02956; AAA33150.1; JOINED. SEQUENCE 11 AA; 1205 MW; 728B4C14C6C2
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Eukaryota; Mycetozoa; Dic
                                                                                                                                                                                                                                                                              NCBI_TaxID=39404;
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Limnodynastes salmini (Salmin's-striped frog).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;

Limnodynastinae; Limnodynastes.
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Last annotation updat
                                                       3AB5A976CAA72728
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annotation
                                                                                             J.H., Wallace J.C., 7 ne structure of the dy in from Limnodynastes
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46:495-505(1982)
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1.1e+04;
                                                                                          Wallace J.C., Tyler M.J.;
cucture of the dynastins from
Limnodynastes fletcheri.";
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RESULT 13
Q40659
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AC Q4065
DT 01-NC
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Q40659
Q40659;
01-NOV-1996
01-NOV-1996
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Q05403;
Q1-NOV-1996
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MEDLINE=84221901; Pubm
Friedman D.J., Umeda P
Frabinowitz M.;
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EMBL; X83121; CAA58183.1;
MON TER 8 8 770 MW.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 22, Last annotation update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
DNA for ORF'S from chromosome XV (Fragment).
Baccharomyces cerevisiae (Baker's yeast).
Eukaryota, Fungi, Ascomycota, Saccharomycotina, Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Alpha-myosin heavy chain (Fragment).
Oryctolagus cuniculus (Rabbit).
Dukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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Q28742;
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STRAIN=FY1679;
MEDLINE=98021609; PubMed=8533473;
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PIR; I46868; I46868.
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Proc. Natl. Acad. Sci. U.S.A
EMBL; K01698; AAA31415.1; -.
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[1]
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eda P.K., Sinha
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U.S.A. 81:3044-3048(1984)
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Pred. No. le+0
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1e+06;
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chromosome XV con reading frames.
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RESULT 15
P82083
ID P8208
AC P8208
DT 01-MAD
DT 01-MAD
DT 01-MAD
DE DYNAS
C Eukar
OC Amphi
OC Limno
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01-MAY-2000
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                    Limnodynastes salmini (Salmin's-striped fi
Eukaryota; Metazoa; Chordata; Craniata; V
Amphibia; Batrachia; Anura; Neobatrachia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.:
"Peptides from Australian frogs. The structure of the dynastins Limmodynastes salmini and Fletcherin from Limmodynastes fletcher Aust. J. Chem. 46:1235-1244(1993).

-i- MASS SPECTROMETRY: MW=772; METHOD=FAB.
SEQUENCE 8 AA; 772 MW; 7BS8772455A2C728 CRC64;
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    Limnodynastinae; Limnodynastes
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NCBI_TaxID=39404;
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Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;
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MEDLINE=91078641; PubM
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Ehrhartoideae; Oryzeae; Oryza.
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Oryza sativa (Rice).
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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h M., Terashima
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1e+06;
frog).
Vertebrata; Euteleostomi;
.a; Bufonoidea; Myobatrach:
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The invention formation and

relates to a new antigenic peptide for inducing the isolation of antibodies having an affinity to it, being

Claim 1; Page 5; 11pp; English.

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prot	HLA prote	HLA prote	PTH(1-14)	N-termina	Human hae	Anti CD34	Antibody	Human DNA		Human par	Antigenic	Human par	PTH-rp N-	Parathyro	PTH(1-14)	Parathyro	Rat parat	Parathyro	Parathyro

ALIGNMENTS

11-FEB-2003

(first

entry)

ABG72607;

Parathyroid hormone antigenic peptide 2-12.

ABG72607 standard; peptide; 11

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RESULT 1
ABG72607
ID ABG72607
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XX ABG7
XX ABG7
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XX Huma
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Rattus sp.
Bos taurus.
Sus scrofa.
                                                                                                                                                                                                                        New parathyroid hormone (PTH) antigenic peptide inducing the formation and isolation of antibodies having an affinity to it, useful for determining bioactive PTH levels in serum, plasma and/or cell culture
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(LAVI/) LAVIGNE J R.
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WPI; 2003-066685/06

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RESULT 2
ABG72608
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC are; (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one Crists peptide antigen comprising a peptide fragment of PTH; (b) CC administering the first peptide antigen to a host animal to induce CC antibody production; (c) monitoring the antibody titre produced; (d) CC isolating antisera produced in the host animal; and (e) selecting CC antisera from the isolated antisera produced in the host that is capable CC of binding to a second peptide antigen; (2) an antibody (ab) produced by the method; and (3) test kits and analytical procedures used for the CC compositions of the present invention are useful for determining CC disactive intact PTH levels in serum, plasma and/or cell culture media. The antigens, antibodies and methods of the present invention, as CC compared to prior art, have the particular advantages of possessing CC greater affinity for FTH, and in particular advantages of possessing CC produced to prior art have negligible cross-reactivity with the large con-molecular forms of PTH. PTH levels are an important parameter in patients suffering from hypercalcaemia, osteoporosis and primary compared to prior art have negligible cross-reactivity with the large con-molecular forms of PTH. PTH levels are an important parameter in patients suffering from hypercalcaemia, osteoporosis and primary compared to primary animo acids 2-12 of human, mouse, rat, porcine, CC canine and bovine PTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 10; Conserv
                               (ZAHR/)
(LAVI/)
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Rattus sp
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Zahradnik RJ
                                                                               05-DEC-2000; 2000US-00730174.
                                                                                                                 05-DEC-2000; 2000US-00730174.
                                                                                                                                                  15-AUG-2002
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is familiaris
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                              ZAHRADNIK R J.
LAVIGNE J R.
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Lavigne JR;
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                                                                                                                                                                                                                                                                                Location/Qualifiers
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Sequence 12 AA;

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CC are; (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one C first peptide antigen comprising a peptide fragment of PTH; (b) CC administering the first peptide antigen to a host animal to induce CC antibody production; (c) monitoring the antibody titre produced; (d) CC isolating antisera produced in the host animal; and (e) electing CC antibody produced antisera produced in the host that is capable CC the method; and (3) test kits and analytical procedures used for the CC the method; and (3) test kits and analytical procedures used for the CC compositions of the present invention are useful for determining CC bioactive intact PTH levels in serum, plasma and/or cell culture media. CC The antigens, antibodies and methods of the present invention, as CC compared to prior art, have the particular advantages of possessing CC greater affinity for PTH, and in particular, are designed to have a novel CC patients suffering from hypercalcaemia, osteoporosis and primary CC hyperparathyroidism. The present sequence represents a PTH antigenic paptide representing amino acids 1-12 of human, mouse, rat, porcine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, being formed from the N-terminus of parathyroid hormone (PTH). Also included
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 5; 11pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New parathyroid hormone (PTH) antigenic peptide inducing the formation and isolation of antibodies having an affinity to it, useful for determining bloactive PTH levels in serum, plasma and/or cell culture
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RESULT 3
AAY96968
AAY96968
AAY96968
AC AAY9
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AC PITH,
KW PITH,
KW bone
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parathyroid hormone N-terminal signaling domain (residues 1-11).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY96968 standard; peptide; 11 AA
    (GARD/) GARDELLA T J.
(KRON/) KRONENBERG H M
(POTT/) POTTS J T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             parathyroid hormone; N-terminal; signaling domain; bone mass, reformation; resorption; remodeling; tether1; osteoporosis.
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Pred. No. 0.014;
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Compounds of the structure or formula S-(L) n-B, R 1-S-(L) n-R or S-(L) n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-TP) (1-34); R 1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTH-P which avoids the need for regular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
             Claim
                                     New antigenic peptide(s) fr
generated using them, able
forms of the hormone.
                                                                                                                                                                                                                                       04-APR-1996
                                                                                                                                                                                                                                                                                                                          diagnosis;
                                                                                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                                                                                 Human parathyroid hormone antigenic peptide hPTH 1-10
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                                                                                              WPI; 1996-180391/19
                                                                                                                                                      (FORS/)
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            2,
                                                                                                                                                                                                                                                                                                                                      parathyroid
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          Page 4; 5pp;
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                                                                                                                                                                                94DE-04434551.
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                                                                                                                                                                                                                                                                                                                          hormone; hPTH; antigen; alpha-helix; hPTH 1-37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide; 10
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             German
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MH
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Pred. No. 0.02;
                                                    human parathyroid hormone - and antibod: distinguish between active and inactive
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                                                                                                                                                                                                                                                                                                                                        immunogen;
                                                                   and antibodies
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Query Match Best Local S Matches 8

Similarity 8; Conserv

Conservative

68.9%;

Score 42; Pred. No.

DB 3;

Length 10

Mismatches 0

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Gaps

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Sequence

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AA

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RESULT 5
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Best Local :
                                                                                   The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipolyhilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin first the thighs and hips, in the treatment of cellulite and for skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a specific example of claimed immunogenic peptides having a sequence from hPTH (1-37) which includes the N- or C-terminal alpha-helical region and/or the non-structured region of the hormone. Antibodies and their binding fragments generated by injecting animal with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                         New parathyroid hormone fragment peptides, in topically applied cosmetic compositions excessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
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Pred. No. 0.37
1; Mismatches
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0.37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             used as lipolysis stimulants for slimming treatment of
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RESULT 6
AAB86219
RESULT 7
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ID ABR4
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AC ABR4
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                                                                                                                                                                                                                                                                                                     This invention describes a novel method for determining (M1) the content (CC of active parathyroid hormone (A) by treating a sample with (1) antibody (CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), (CC and including the N-terminal residue and (ii) antibody (Ab2) that (CC recognizes an epicope within the receptor-binding site of (A). The number (CC of molecules that react with both antibodies is determined and used to (CC calculate the content of physiologically active (A). The method is used (A) reparathyroidism. The method (importers, osteopathy, or hypo- or (CC of parathyroidism. The method (impike known assays) recognizes that (CC (i) some fragments of (A) shorter than the complete (B4 aa) peptide are active and (ii) that apparently intact peptide may be biologically (CC care antagonistic (these have the receptor-binding site but lack the N-CC (CC terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This (CC sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                              Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
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                   ABR44166;
                                                ABR44166 standard; peptide; 10
                                                                                                                                                                                                                                                             Sequence 10 AA;
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                                                                                                                                                                                              Similarity 80.0
8; Conservative
                                                                                                                                 SVSEIQLMHN
                                                                                                                                                             AVSEIGFMHN 10
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                                                                                                                                   10
                                                                                                                                                                                                               68.9%;
80.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunogenic peptide SEQ ID 1
                                                                                                                                                                                              Score 42; DB 4;
Pred. No. 0.37;
1; Mismatches
                                                   ĀĀ
                                                                                                                                                                                                                             Length 10;
                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                 Gaps
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RAESULT 8
AAYSOGN
ID AAAYS
XX AAYS
XX AAYS
XX AAYS
XX Cyc.
XX 
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Song Y,
Chang M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a fusion peptide (Tat-hPTHDP), where self cell-penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmetic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endothelium, does not cause skin disease and has superior lipolysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 5; Page 6; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel fusion peptide comprising self to human parathyroid hormone-derived slimming cosmetic composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fusion peptide; tat; hPTHDP; parathyroid hormone; skin; cosmetic;
lipolysis; human; hPTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-468288/44.
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15-MAR-2002; 2002KR-00014062.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2003
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                                                                                                                                                                                               Cyclic peptide; resin bound; parathyroid hormone; osteopathic; disea; hypotensive; PTH receptor; treatment; hyper-calcemia; hypo-calcemia; osteoporosis; osteopenia; hyper-parathyroidism; hypo-parathyroidism; Cushing's syndrome; renal failure; hypertension; bone fracture repai:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY50600 standard; peptide; 11 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GLDS )
                          Misc-difference
                                                                                                                                                                                                                                                                                                                                Resin bound
                                                                                                                                                                                                                                                                                                                                                                                09-FEB-2000
                                                                           Misc-difference
                                                                                                                                                      Synthetic
                                                                                                                                                                                            Cushing's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    parathyroid hormone (hPTH) fragment (residues 1-10).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LG HOUSEHOLD & HEALTH CARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kang N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SVSEIQLMHN 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AVSEIOFMHN
                                                                                                                                                                                                                                                                                                                                cyclic peptide 33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002WO-KR000835
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                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                              Location/Qualifiers
/note= "Ser(OtBu)"
                                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Park S,
                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cho W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 42; DB 6;
Pred. No. 0.37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cell-penetrating peptide, useful a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ູເດ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lee Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 10
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as component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                  disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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of skin
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Misc-difference

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ARBSULT 9
AAB86225
ID AAB8
XX AAB8
XX AAB8
XX Huma
XX Para
XX Huma
                                                                                                                                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for the preparation of peptides (II) that contain both cyclic and linear peptide fragments comprises consequential reaction of a resin-bound linear fragment with the cyclic fragment in N-protected form and optionally other linear fragments. The products of the invention have osteopathic and hypotensive activity. (II) think the cyclic method is particularly used to prepare cyclic peptide analogs of parathyroid hormone (PTH) or PTH-related peptides which are useful for treating diseases that respond to treatment with agents that bind to PTH creeptors (with or without activation of adenyl) cyclase activity). e.g. hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-garathyroidism, Cushing's syndrome, renal failure and hypertension, also for promoting repair of bone fractures. Separate synthesis of the cyclic fragment allows convergent synthesis of resin-bound (II), with better cyields and higher throughput. The difficulties associated with cyields and higher throughput. The difficulties associated with cyields and higher throughput. The difficulties associated with the peptide before reaction with the resin-bound component. AAY50568-creation of the bridged fragments are confined to a small peptide which can be purified before reaction with the resin-bound component. AAY50568-creation of the bridged fragments described in the method of the
                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                 Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                               Human parathyroid
                                                                                                                                    03-SEP-2001
                                                                                                                                                                    AAB86225;
                                                                                                                                                                                                    AAB86225 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 75; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
                               hypo-parathyroidism;
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RHON ) RHONE-POULENC RORER PHARM INC
                                                                                                                                                                                                                                                                                                                                                   10;
                                                                                                                                                                                                                                                                                                          1 AVSEIQFMHNLG 12
                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AW,
                                                                                                                                                                                                                                                                                   AVSEIQ-LHNLG 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          synthesis of peptides for treating e.g. bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                                      ĀΑ;
                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mencel
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                                                                                                  hormone
                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                    68.0%;
                               hyper-parathyroidism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Asn(Trt)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "His(Trt)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Glu (OtBu)
                                                                                                 immunogenic peptide SEQ
                                                                                                                                                                                                                                                                                                                                                                    Score 41.5;
Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                      ed. No. 0.5;
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                    Length 11;
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RESULT 10
AAB01862
ID AAB01
XX AAB01
XX I1-SE
XX Parat
KW Calci
KW Calci
KW Calci
KW Calci
XX Homo
OS Synth
XX WO200
XX Y 27-AF
XX 20-OC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC (Ab1) that recognizes the epitope formed by amino acids (a) 1-3 of (A).

CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A).

CC and including the N-terminal residue and (ii) antibody (Ab2) that

CC enoughizes an epitope within the receptor-binding site of (A). The number

CC of molecules that react with both antibodies is determined and used to

CC calculate the content of physiologically active (A). The method is used

CC of riagnosis of calcium-metabolism disorders, osteopathy, or thypo- or

CC typer-parathyroidism. The method (unlike known assays) recognizes that

CC (i) some fragments of (A) shorter than the complete (B4 aa) peptide are

CC active and (ii) that apparently intact peptide may be biologically

CC are antagonistic (these have the receptor-binding site but lack the N-

CC terminus). It thus provides a true measure of the content of

CC and its 1-37 fragment which may produce falsely high values. This

CC sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                           Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthes bone synthesis; agonist; osteoporosis; non-parenteral deliver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibod reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 3; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-376318/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-DEC-1999;
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                                                                                                       Synthetic
                                                                                                                          Homo sapiens
                                                                                                                                                                                                                               PTH(1-14)/PTHrP(1-14)-derived peptide,
                                                                                                                                                                                                                                                                  11-SEP-2000
                                                                                                                                                                                                                                                                                                      AAB01862;
                                                                                                                                                                                                                                                                                                                                     AAB01862 standard; peptide; 9
                                                                      WO200023594-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                          1 VSEIQLMHN 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 VSEIQFMHN 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 41; DB 4; Li
Pred. No. 1.4e+06;
0; Mismatches 1;
                                                                                                                                                                                                                               SEQ ID
                                                                                                                                                                                                                                 NO: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          two antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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20-OCT-1999; 27-APR-2000.

99WO-US024481

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RESULT 11
AAY96966
ID AAY96
XX
AC AAY96
XX
DT 31-OC
XX
EW PTH;
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                                                                                                                                                                                                                                                                                                                                                                                                                             CC BO1869). The peptides of the invention are at least 85 identical to the CG Gly-Lys-X6 (AAB01857) where: X1 is Ser of Ala; X2 is ILe or Lys; X3 is CC Met, Leu or NIe; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; CC provided that the peptide is not PTHIP(1-14). The peptides of the CC invention also encompass fragments of peptides of the invention control of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-CC terminal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal, cc skeletal, neurological system, neuromuscular and cardiovascular systems. CC it binds to both PTH-1 receptors on osteoblasts and renal tubular cells, and to the recently identified PTH-2 receptor. PTH has a potent anabolic calls of protein hormones, the PTH-2 receptor. PTH has a potent anabolic calls of protein hormones, the PTH-2 receptor. PTH principles of the renal and skeletal actions of PTH, and also bind to the PTH-1 calcium reabsorption, enhances calls of protein bind to the PTH-2 receptor. The peptides of the cenal and skeletal actions of PTH, and also bind to the PTH-1 calcium reabsorption. PTH-1 PTH-2 receptor antagonists are useful for the treating conditions characterised by a decrease in bone mass, such as calcium schement of the PTH-1 receptor. Detectably labelled peptides of the invention are also useful in the determination of rates of bone formation, bone resorption conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via convention are peptides.
                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Farathyroid hormone (PTH) peptides, acids that encode them, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel parathyroid hormone (PTH) (AAB01859) and parathyroid hormone-related peptide (PTHrP, biologically active derivatives thereof (AAB01857-B01858, i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KRON/)
                                  Parathyroid hormone N-terminal signaling domain (residues 1-9).
                                                                       31-OCT-2000
                                                                                                          AAY96966
                                                                                                                                             AAY96966 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GARD/) GARDELLA T
                                                                                                                                                                                                                                                                                                            Similarity
8; Conserv
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POTTS J T.
                                                                                                                                                                                                                                                                       AVSEIQFMH 9
                                                                                                                                                                                                                                        AVSEIQLMH
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                                                                                                                                                                                                                                                                                                                                                                                                                     receptor agonists
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                                                                       (first
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 hormone;
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                                                                     entry)
                                                                                                                                                                                                                                                                                                                                63.9%;
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 N-terminal;
                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                Score 39; DB 3;
Pred. No. 1.4e+06;
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                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating osteoporosis
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 signaling
                                                                                                                                                                                                                                                                                                                                                 DB 3;
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                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                 Length 9;
 domain;
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 bone mass
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compounds of the structure or formula S-(L) n-B, R 1-S-(L) n-R or S-(L) n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R 1 is the PTH-1 receptor signal sequence; and R is (a portion of) pTH-1 receptor sequence; and R is (a portion of) pTH-1 receptor sequence of bone removed by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalaction unlike the large native PTH or PTH PTH which avoids the need for regular injections to treat osteoporosis
                                                                                                                                                                                                                      Human parathyroid hormone; hPTH; antigen; alpha-helix; immunogen; diagnosis; active hPTH 1-37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
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(KRON/) KRONENBERG
(POTT/) POTTS J T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (JUEP/)
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) POTTS J T.
) JUEPPNER H.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; peptide; 9
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                                                                                                                                                                                                                                                                                                              hormone antigenic peptide hPTH 1-9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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Pred. No. 1.4e+06;
0; Mismatches 1
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04-APR-1996

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28-SEP-1994;

94DE-04434551

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RESULT 13
AAB01863
ID AAB01
XX AAB018
AC AAB01
AC AAB01
XX AAB01
AC AAB01
XX AAB01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 77.8 Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                              Parathyroid hormone peptide; PTH; PTH-related peptide; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D bone synthesis; agonist; osteoporosis; non-parenteral c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a specific example of claimed immunogenic peptides having a sequence from hFTH(1-37) which includes the N-terminal alpha-helical region and/or the non-structured region of hormone. Antibodies and their binding fragments generated by injanimal with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antigenic peptide(s) from generated using them, able to forms of the hormone.
                                                Parathyroid hormone (PT) acids that encode them,
                         Disclosure; Page 26;
                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB01863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB01863 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9 AA;
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                                                                                                                Gardella TJ,
                                                                                                                                                      (KRON/)
                                                                                                                                                                                                        22-OCT-1998;
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                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:7
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                                                                                         2000-339693/29
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                                                                                                                                                    GARDELLA T J.
KRONENBERG H M.
POTTS J T.
                                                                                                                                        JUEPPNER H.
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                                                                                                              Kronenberg
                                                                                                                                                                                                        98US-0105530P
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77.8%;
                        73pp; English
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                                                  H) peptides,
useful for
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Pred. No. 1.4e+0
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                                                                                                                Potts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                treating
                                                                                                              JT,
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                                                              PTH related peptides and the nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .4e+06;
                                                                                                              Jueppner
                                                  osteoporosis
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delivery.
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novel parathyroid

hormone

(PTH) peptide

Gardella

IJ,

Kronenberg

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Potts ŢŢ,

Jueppner

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RESULT 14
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XX PTH;
XW bone
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Best Local
                   (GARD/)
(KRON/)
(POTT/)
                                                                                                                                                                        AAY96981 standard; peptide;
                                                                                                                                                                                                                                                             Sequence 9 AA;
                                              31-DEC-1998;
                                                          30-DEC-1999;
                                                                       06-JUL-2000.
                                                                                     WO200039278-A2
                                                                                                                                  Parathyroid hormone N-terminal signaling domain.
                                                                                                                                               31-OCT-2000
                                                                                                                                                             AAY96981;
                                                                                                 sapiens
                                                                                                              parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tetherl; osteoporosis.
            GARDELLA T J.
) KRONENBERG H I
) POTTS J T.
) JUEPPNER H.
                                                                                                                                                                                                                                   Similarity 7; Conserv
                                                                                                                                                                                                                      AVSEIQFMH
                                                                                                                                                                                                          AVSEIQLLH
                                                                                                                                                                                                                                    Conservative
                                                                                                                                               (first entry)
                                              98US-0114577P
                                                           99WO-US031108
                                                                                                                                                                                                                                          59.0%;
77.8%;
                                                                                                                                                                                                                                   Score 36; DB : Pred. No. 1.4e.
1; Mismatches
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1.4e+06;
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RESULT 15
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-TP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTH-PTH which avoids the need for regular injections to treat osteoporosis
                                                                             Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                               WPI; 2001-376318/40
                                                                                                                                                                            Armbruster FP
                                                                                                                                                                                                             (-DWMI)
                                                                                                                                                                                                                                              17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                      Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human parathyroid hormone immunogenic peptide SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
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77.8%;
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This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody

Disclosure; Page 3; 10pp; German

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                                                                                                                                                                                                                                          CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CC for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or CC hyper-parathyroidism. The method (unlike known assays) recognizes that CC (i) some fragments of (A) shorter than the complete (84 aa) peptide are CC active and (ii) that apparently intact peptide may be biologically CC inactive, and also takes into account the fact that some fragments of (A) creminus). It thus provides a true measure of the content of CC terminus). It thus provides a true measure of the content of CC and its 1-37 fragment which may produce falsely high values. This CC sequence represents a peptide fragment used to illustrate the method of the invertion.
                                                                                    Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                   Sequence 9 AA;
                                                                                                                                                                                                                                 invention
SVSEIQLMH
                                              AVSEIQFMH 9
                                                                                         Conservative
  ω
                                                                                                               59.0%;
77.8%;
                                                                                            1; Mismatches
                                                                                                               Score 36; DB 4;
Pred. No. 1.4e+06;
                                                                                                                                     Length 9;
                                                                                            Indels
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Search completed: May 18, Job time : 45.75 secs 2004, 10:01:54

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                Score
                Query
Match
             US-09-730-174A-6
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100.425 Million cell updates/sec
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Gapop 10.0 , Gapext
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/cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*
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                                                                                                                                                                                                                                                                                                Length
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         US-09-730-174A-6
US-09-730-174A-4
US-09-730-174A-5
US-09-730-174A-5
US-09-730-174A-1
US-09-730-174A-1
US-09-730-174A-1
US-10-168-185-7
US-10-168-185-7
US-10-168-185-7
US-10-168-185-8
US-10-188-185-8
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Sequence 6, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 7, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 7, Appli
Sequence 8, Appli
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RESULT 1 US-09-730-174A-6

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16
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equence 26, App	equence 508, Ap	equence 4, Appl	equence 5, Appl	mence 16,	equence 14, App	equence 61, App	equence 26,	quence 78,	equence 73,	equence 55,	equence 45, Appl	ence 13	equence 12, App	equence 3, Appl	equence 526, Ap	equence 115, Ap	equence 49	equence 18, App	equence 12	equence 108, Ap	equence 12	equence 19, App	equence 4.,	equence 41, App	quence 12,	quence 4,	equence 4, Appli	ence 10, Ap	equence 2, A

ALIGNMENTS

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RESULT 2
US-09-730-174A-4
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Patent No. US2002010871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having of INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hole REFERENCE: IMUNE-001A
Sequence 4, Application US/09730174A

Patent No. US20020110871A1

GENERAL INFORMATION:

APPLICANT: Zahradnik, R.J.

TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid HC

FILE REFERENCE: INUNE-001A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 6
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Best Local (
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ORGANISM: Artificial Sequence
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12; Conserv
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RESULT 4
US-09-730-174A-5
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US-09-730-174A-2
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Best Local S
Matches 11
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/09730174A Patent No. US20020110871A1 GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid FILE REFERENCE: INUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09730174A Patent No. US20020110871A1 GENERAL INFORMATION:
                                                                                                                                                                           SEQ ID NO 5
LENGTH: 12
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 4
LENGTH: 12
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Induces Formation of Antibodies and Isolates said Antibodies -09-730-174\text{A}-2
                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid He FILE REFERENCE: IMUNE-001A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zahradnik, R.J
                                                                                             FEATURE:
OTHER INFORMATION: Induces Formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
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                                        Local
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                      Conservative
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                                     90.2%;
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91.7%;
                    Score 55; DB 9;
Pred. No. 0.0015;
0; Mismatches
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Pred. No. 0.0006;
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Pred. No. 0.00043;
1; Mismatches (
                                                                                                           of Antibodies
                                                    Length 12;
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                                                                                                                                                                                                                                                                                   Matches
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Best Local :
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
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RESULT 7
US-10-168-185-1
                                                                                  Sequence 1, Application US/10168185; Publication No. US20030175802A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho: FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
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Best Local S
Matches 10
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Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho:
TITLE REFERENCE: INUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NO3
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US Patent No. US20020110871A1
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidd-Gayk, Heinric
APPLICANT: Roth, Heinz-Jurgen
                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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10; Conser
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  Missbichler, Albert
Schmidt-Gayk, Heinrich
Roth, Heinz-Jurgen
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Pred. No. 0.0078;
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Pred. No. 0.0055;
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PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR PILING DATE: 1999-12-17
NUMBER OF SEO ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
US-10-168-185-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/ED00/12911
PRIOR APPLICATION NUMBER: PCT/ED00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR PILING DATE: 1999-12-17
PRIOR FILING DATE: 1999-12-17
NUMBER: OF SEQ ID NOS: 11
SOFTWARE: FRSESEQ for Windows Version 4.0
SEQ ID NO 7
                                                                                                                                                                                                                                                   RESULT 9
US-10-192-673-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens US-10-168-185-7
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Broactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570002
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Best Local S
Matches 8
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Publication No. US20030175802A1
GENERAL INFORMATION:
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Best Local (
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Juzgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
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CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
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TYPE: PRT
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les 8; Conser
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Local Similarity 80.0%;
hes 8; Conservative
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Pred. No. 0.33;
1; Mismatches
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APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEG ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 6
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US-10-192-673-7
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Matches
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Publication No. US20030166838A1
GENERAL INFORMATION:
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Publication No. US20030175802A1
GENERAL INFORMATION:
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potte, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTHr)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Uurgen
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TYPE: PRT
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TYPE: PRT
ORGANISM: Artificial Sequence
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87.5%;
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Pred. No. 1e+0
0; Mismatches
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Pred. No. 1e+06;
0; Mismatches
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Reth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
TITLE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: DCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
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Best Local Similarity
7; Conserve
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                                                                                                                                                                                               US-10-192-673-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo sapiens
US-10-168-185-2
                                                         Sequence 8, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
7; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
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CURRENT FILLING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Armbruster, Franz Paul
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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ORGANISM: Artificial Sequence
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77.8%;
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Pred. No.
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RESULT 15 US-10-232-187-8

GENERAL INFORMATION:

APPLICANT: Abrahamson, APPLICANT: Bochner, B

Julie A.

APPLICANT: APPLICANT: APPLICANT:

Bochner, Bruce Erickson-Miller, Connie L. Kikly, Kristine K. Schleimer, Robert Sequence 8, Application US/10232187 Publication No. US20030092091A1

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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/10168185 Publication No. US20030175802A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                               SEQ ID NO 3
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Best Local (
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                                                                              Best Local Similarity
Matches 6; Conserv
                                                                                                                       Query Match
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT APPLICATION NUMBER: US/10/168,185
                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER: FILING DATE: 1999-12-17
                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Hormone-Related Peptide (FTHrP) FILE REFERENCE: 0609.4570002
                                                                                                                                                                           TYPE: PRT ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                     1 AVSEIQFM 8
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SVSEIQLM
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75.0%;
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Pred. No. 1e+06;
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                                                                                Mismatches
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APPLICANT: Nulku, Turkan E.

TITLE OF INVENTION: Sialoadhesin Factor-2 Antibodies

FILE REFERENCE: GH50042-1

CURRENT APPLICATION NUMBER: US/10/232,187

CURRENT FILING DATE: 2002-08-29

FRIOR APPLICATION NUMBER: 60/187,955

PRIOR APPLICATION NUMBER: 60/315,943

PRIOR FILING DATE: 2001-03-05

PRIOR FILING DATE: 2002-01-18

PRIOR FILING DATE: 2002-07-10

PRIOR FILING DATE: 2002-07-10

VUMBER OF EXC ID NOS: 14

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 8

LENGTH: 10

TYPE: PRT

ORGANISM: Homo Sapien

US-10-232-187-8

Ouery Match

Best Local Similarity 44.4%; Score 27; DB 14; Length 10;

Best Local Similarity 44.4%; Pred. No. 2e+02;

Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps

Oy

1 AVSETOFMH 9

Db 2 ATSSVSYMH 10

Search completed: May 18, 2004, 10:21:00

Search completed: May 18, 2004, 10:21:00
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Result
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Maximum DB
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Listing first 45 summaries
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Match
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                                                                                                                                                                                                                                                                                                                                                               is the number of results predicted by chance to have a ster than or equal to the score of the result being printed, rived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                      Issued_patents_AA:*
1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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   AVSEIQFMHNLG
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Copyright (c) 1993 - 2004 Compugen Ltd.
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 US-09-442-989-31
US-09-421-379-6
US-09-421-379-7
S460978-3
US-09-421-379-8
US-09-421-379-8
US-09-421-379-10
US-08-1817-547A-3
US-08-482-528-42
US-08-482-528-42
US-08-483-528-42
US-08-483-562-9
US-08-483-562-9
US-08-483-561-9
US-08-483-561-9
US-08-483-561-9
US-08-12-378-12
US-09-391-3858-108
US-09-391-371-2
US-09-391-371-3
US-08-687-706-56
US-08-817-547A-15
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Sequence 1, Appli
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Sequence 9, Appli
Sequence 9, Appli
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Sequence 12, Appli
Sequence 13, Appli
Sequence 14, Appli
Sequence 56, Appli
Sequence 16, Appli
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US-09-442-989-31
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ALIGNMENTS

		-989-31
	Nie	ORMATION:
		IOCATION: (8)
		NAME/KEY MOD RES
	Asn(Trt)	OTHER INFORMATION:
		FEATURE:
	His(Trt)	0
		NAME/KEY: PEPTIDE
		FEATURE:
	Gln (Trt)	o .
		LOCATION: (6)
		•••
		FEATURE:
	Glu (OtBu)	O.
	-	
		NAME/KEY: PEPTIDE
		FEATURE:
	Ser (OtBu)	FO
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		FEATURE:
	FMOC-Ala	О
		FEATURE:
	1 Sequence	ORGANISM: Artificial
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	: 46	O
	FILING DATE: 1998-04-15	
	NUMBER: 60/081,897	
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	NUMBER: US/09/442,989	
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RESIN-BOUND CYCLIC	ESS FOR THE PREPARATION OF	NVENTION
	ч	APPLICANT: Mencel, J
	Adam W.	APPLICANT: Sledeski,
		Z.
	ion IIS/09442989	Semience 31 Application
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DB 4;

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Query Match
Best Local Similarity
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US-08-817-547A-1
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                                                                                                                      RESULT 3
US-09-421-379-6
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: no
Sequence 6, Application US/09421379
Patent No. 6495662
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08817547A Patent No. 6030790
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Flopy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentth Release #1.0, Vers
SOFTWARE: Patentth Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
APTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REGISTRATION NUMBER: 22,176
REGISTRATION INFORMATION:
TELEPHONE: 404-818-3700
TELEPHONE: 404-818-3700
TELEPAX: 404-818-3790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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APPLICANT: F
APPLICANT: N
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of COMPUTER: IBM FC comp
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TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U
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STATE: Georgia
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191 Peachtree Street, 37th Floor
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Hock, Dieter
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Pred. No.
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                                                                                                                                                                                                                                                                            Score 42; 1
Pred. No. 0
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RESULT 4
US-08-817-547A-2
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LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PATENT NO. BURNETION:
GAPPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone Related Peptide (PTHrP) FILE REFERENCE: 0609.4570001 CURRENT APPLICATION NUMBER: US/09/421,379 CURRENT FILING DATE: 1999-10-20 EARLIER APPLICATION NUMBER: US/09/421,379 EARLIER FILING DATE: 1998-10-20 EARLIER FILING DATE: 1998-10-22 NUMBER: US/09/530 EARLIER FILING DATE: 1998-10-22 NUMBER OF SEQ ID NOS: 13
                                                                                                           TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                 PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION UNMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
TOPULCE: pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Atlanta
STATE: Georgia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE:
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                                                     TYPE: amino acid
STRANDEDNESS: unl
                                                                                                                                                                     TELEPHONE:
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8; Conserv
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                                                                                             9 amino acids
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                                         unknown
                                                                                                                                                                       404-818-3700
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                                                         unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63.9%;
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Pred. No. 3e+05;
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; HYPOTHETICAL: ; ANTI-SENSE: 1 US-08-817-547A-2

peptide

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Query Match Best Local

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PATERIO NO. 5460978

APPLICANT: MARTIN, THOMAS J.; MOSELEY, JANE M.; KEMP,

BRUCE E.; WETTENHALL, RICHARD E.H.

TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL

; PPERCALCEMIA OF MALIGNANCY-PFHRP

NUMBER OF SEQUENCES: 4

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/715,280

FILING DATE: 14-JUN-191

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 199,235

FILING DATE: 09-MAY-1988

APPLICATION NUMBER: 199,235

FILING DATE: 09-MAY-1988

APPLICATION NUMBER:
                                       Query Match
Best Local Similarity
7; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: synthetic ; OTHER INFORMATION: peptide US-09-421-379-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
ETITE DEFENDED. ACCO. ACCO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 7
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Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUWBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity nes 7; Conservat
                                                                                                                                                                 LENGTH: 11
                                                                                                                                                                                                        FILING DATE:
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AVSEIQFMHN 10
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                                                            Conservative
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77.8%;
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77.8%;
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70.0%;
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Pred. No. 3e+05;
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                                                                                Score 31; DB
Pred. No. 8.5;
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APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Branthyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION UNUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER FILING DATE: 1998-10-22
NUMBER OF 520 ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SBQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-8
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US-08-140-137A-42
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US-09-421-379-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 42, Application US/08140137A
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMÁTION:
APPLICANT: TUDMANEN, ELAINE
APPLICANT: MASURE, H. R.

PITILE OF INVENTION: ANALOGS OF ENDOTHELIAL LEUKOCYTE
TITLE OF INVENTION: ADHESION MOLECULE (ELAM)
NUMBER OF SEQUENCES:
ADDRESSEB: Klauber & Jackson
ADDRESSEB: Klauber & Jackson
                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION TOWNBER: US/08/140,137A
FILING DATE: 27-MAY-1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esg., David A.
REGISTRATION NUMBER: 26,742
REGISTRATION NUMBER: 26,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                   REGISTRATION NUMBER: 26,742
REFERENCE/POCKET NUMBER: 600
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-5800
                                                                                                                                                                                                                                                                                                                                                                                          STREET:
CITY: H
                                                                                                                                                                                                                                                                                                                                07601
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: New Jersey
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  201 343-1684
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                                                            600-1-096
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Pred. No. 3e+05;
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                                                                                                                                                                                                                              Version #1.25
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RESULT 6 5460978-3

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Matches Query Match Best Local (

ENGTH:

5460978-3

SEQ ID NO:3:

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TOPOLOGY: linear
HOLECULE TYPE: peptide
DESCRIPTION: CaRD peptide sequence
US-08-140-137A-42
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US-08-817-547A-3
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Best Local Similarity 62.5%;
Matches 5; Conservative
                                                             Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: PROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-18-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION: APPLICANT: Aderma
                                                                                                                                                                                                                                                  TELEPHONE: 404-818-3700
TELEPAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                              TOPULUE TYPE: pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Jones & A
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APPLICANT: Magerlein, M
                                                                                                                                               HYPOTHETICAL:
ANTI-SENSE: I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 191 Peachtree Street, 37th Floor
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
                                                                                                                                                                                                    TOPOLOGY: unl
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5. 6030790
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                              1 AVSEIQFM 8
                                                                                                                                                                                                                                   amino acid
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                                                                 Conservative
                                                                                                                                                   no
                                                                                                                                                                                                      unknown
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                                                                                                                                                                                                                     unknown
                                                                                                                                                                                  peptide
                                                                                45.9%;
75.0%;
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Pred. No.
                                                               Score 28; DB 3; Length 8; Pred. No. 3e+05; 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                           07826-0007
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US-08-482-228-42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 6017719
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEAR
NUMBER OF SEQUENCES: 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Guthrie, Janice REGISTRATION NUMBER: 35,170 REFERENCE/DOCKET NUMBER: IT TELECOMMUNICATION INFORMATION: TELEPHONE: (714) 440-5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                     APPLICANT: Al-Abdaly, Fahad A.
APPLICANT: Guillermo, Roy
APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
NUMBER OF SEQUENCES: 215
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
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                                 ADDRESSEE: Janice Guthrie, Ph.D. STREET: P.O. Box 15210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
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GY: linear
Irvine
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                                                                                                                                                                                                                                                                                                                                                                                                                               Tseng-Law, Janet
Kobori, Joan A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Al-Abdaly, Fahad A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44.3%; Score 27; DB 2; 55.6%; Pred. No. 54; tive 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35,170
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STATE: C

California

USA

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GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: Forts, John T.
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
FUNREMY APPLICATION NUMBER: US/09/421,379
CURREMY APPLICATION NUMBER: US/09/421,379
CURREMT FILING DATE: 1998-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
COLTMADE: DEFORTE NOS: 13
COLTMADE: DEFORTE NOS: 13
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                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-10
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                                                              Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 55.6%; Matches 5; Conservative
                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 9
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 92713-5210

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/482,528

FILING DATE: 07-JUN-1995

CLASSIFICATION: 435

CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (714) 553-1952
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (714) 440-5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Guthrie, Janice REGISTRATION NUMBER: 35, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AVSEIQFMH 9
                                     AVSEIQFMH 9
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SVSEHQLLH 9
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                                                                                  Conservative
                                                                                                  55.6%
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Pred. No. 54;
                                                                                                    Score 26; DB 4;
Pred. No. 3e+05;
                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 3;
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                                                                                                                      Length 9;
                                                                                  Indels
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US-08-438-562-9
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US-08-116-778E-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 4
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APPLICANT: NAKAMUR
APPLICANT: KOIKE
APPLICANT: SHITARA
APPLICANT: HANAI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPARTIBLE
COMPUTER: IBM PC COMPARTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.25
SOFTWARE: PACENTIN DATA:
APPLICATION NUMBER: US/08/116,778E
FILING DATE: 07-SEP-93
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
NAME: WILSON, MARY J.
NAME: WILSON, MARY J.
OPERATOR OF THE PROPERTY OF THE PACENT NUMBER ASSETS OF THE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9, Applica Patent No. 5830470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9,
Patent No. !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (703)816-4000
TELEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                   APPLICANT: HASEGAWA, MANORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: KOIKE, MASAMICHI
APPLICANT: SHITARA, KENYA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
ZIP: 22201-4714
STATE: VIRGINIA
COUNTRY: U.S.A.
ZIP: 22201-4714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                    CITY:
                                                                                                                            STREET:
                                                                                                                                                                  ADDRESSEE:
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                                                                                             ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08438562
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                                                                                                                                   : NIXON & VANDERHYE P.C
1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAKAMURA, KAZUYASU
KOIKE, MASAMICHI
                                                                                                                                                                                                                                                                                                                          SHITARA, KENYA
HANAI, NOBUO
KUWANA, YOSHIHISA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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44.4%; Pred. No. 1.1e+02;
ative 2; Mismatches 3;
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US-08-483-528B-97
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             APPLICATION NUMBER: US/08/483,528B
FILING DATE: 07-JUN-95
CLASSIFICATION: 536
TELECOMUNICATION: INFORMATION:
TELEPHONE: (703)816-4100
TELEPACE: (703)816-4100
INFORMATION FOR SEQ ID NO: 97:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 97, Application Patent No. 5939532
Patent No. 5939532
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 41.0%;
Best Local Similarity 44.4%;
Matches 4; Conservative
                                                                                                                                                                                             ADDERSSE: UIXON & VANDERHYE P.C.
ADTRESSE: UIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VIRGINIA
COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPARTINE
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNIMFER - UIS/08/483.5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U5/08/438,562
FILING DATE: 10-MAY-95
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/116,778
FILING DATE: 07-SEP-93
CLASSIFICATION: 424
APPLICATION NUMBER: 03/116,778
FILING DATE: 07-SEP-93
CLASSIFICATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 249-76
TELEPHONE: (703)816-4000
TELEPHONE: (703)816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: HASEGAWA, MITTLE OF INVENTION: HUINUMBER OF SEQUENCES: 1: CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 10 amino acids
TYPE: amino acid
TOPOLOGY: linear
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amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              f: HANAI, NOBUO
f: KUWANA, YOSHIHISA
f: HASEGAWA, MAMORU
INVENTION: HUMANIZED ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08483528B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAKAMURA, KAZUYASU
KOIKE, MASAMICHI
SHITARA, KENYA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   103
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Pred. No. 1.1e+02;
2; Mismatches 3
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; TOPOLOGY: linear;
; MOLECULE TYPE: peptide
US-08-483-528B-97

Query Match
Best Local Similarity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

Popularity 44.4%; Pred. No. 1.1e+02;

Popularity 44.4%; Pred. No. 1.1e+02;

Popularity 44.4%; Pred. No. 1.1e+02;

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